

**RESULT 15**  
**AF465211/c**  
**LOCUS** AF465211 **1768 bp** **DNA** **circular** **VRL 07-FEB-2002**  
**DEFINITION** Porcine circovirus type 2 strain SC, complete genome.  
**ACCESSION** AF465211 **GI:18448942**  
**VERSION** AF465211.1 **GI:18448942**  
**KEYWORDS**  
**SOURCE** Porcine circovirus type 2.  
**ORGANISM** porcine circovirus type 2.  
**REFERENCE** 1 (bases 1 to 1768)  
**AUTHORS** Wang, C., Pan, C.H., Huang, C.C., Huang, T.S., Jong, M.H., Lin, S.Y. and Lai, S.S.  
**TITLE** Complete nucleotide sequences of porcine circovirus type 2 isolated in pigs with various clinical syndromes  
**JOURNAL** Unpublished  
**REFERENCE** 2 (bases 1 to 1768)  
**AUTHORS** Wang, C., Pan, C.H., Huang, C.C., Huang, T.S., Jong, M.H., Lin, S.Y. and Lai, S.S.  
**TITLE** Direct Submission  
**JOURNAL** Submitted (02-JAN-2002) Hog Cholera Department, National Institute for Animal Health, 376 Chung-cheng Road, Tamsui, Taipei 25101, Taiwan  
**FEATURES**  
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**BASE COUNT** 456 **a** 357 **c** 491 **g** 464 **t**  
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Matches 653; Conservative 0; Indels 0; Gaps 0;  
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QY 61 CAGATCCTCCGGCGCCCTGGCTCGTCCACCCGCCACCCGTTACCGCTGGAAAGG 120  
Db 1675 CAGATCCTCCGGCGCCCTGGCTCCACCCGCCACCCGTTACCGCTGGAAAGG 1616  
QY 121 AAAAATGGCATCTCAACACCCGCCTCTCCGCACCTTCGGATATACTGTCAAGCGAAC 180  
Db 1615 AAAAATGGCATCTCAACACCCGCCTCTCCGCACCTTCGGATATACTGTCAAGCGTAC 1556  
QY 181 ACAGTCAGAACGCCCTCCCTGGGGGACATGTGATTCAATATAATGACTTTCTT 240  
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QY 241 CCCCGAGGGGGTCAAACCCCGCTCTGTGCCCTTGAATAAGAAAG 300  
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QY 301 GTTAAGGTGAATTCTGGCCCTGCTCCCCGATCACCAAGGGTACAGGGAGTGGGCTC 360  
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QY 361 AGTCTGTTATTAGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGACCC 420  
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QY 421 TATGTAAACTACTCCGGCATACCAAAACCCGCTTCTCTACCAACTCCGGTAC 480  
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QY 481 TTTACCCCCAACCTGTCCCTAGATTTCACATTGATTACTTCCAAACCAAAAGA 540  
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QY 541 AACCAAGCTGTGGCTGAGACTACAAACTGCTGGAAATGTAGACCACTGGCACT 600  
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Search completed: May 18, 2003, 11:41:48  
Job time : 2030 secs



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Db	197	AACCAGCTGGCTGAGACTCAAACACTGCTGGAAATGTAGACCACTGGGACT	138	
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SOURCE				Db 497 CCCCCAGGGGGCTAAACCCCCGCTCTGTGCCCTTGAATACTACAGATAAGAAAG 438
ORGANISM				QY 301 GTTAAGGGTTGAATTCTGGCCCTGCTCCGCACCCAGGGTACAGGGCCTCACCTATGACCC 360
REFERENCES				Db 437 GTTAAGGGTTGAATTCTGGCCCTGCTCCGCACCCAGGGTACAGGGCCTCACCTATGACCC 378
AUTHORS	Cui, S.J., Li, Y., Li, J.W., Jin, H. and Tong, G.Z.			QY 361 AGTGCTGTATTAGTATACTTGTAAACAAAGGCCACAGCCTCACCTATGACCC 420
TITLE	Isolation and Identification of Porcine Circovirus Type 2			Db 377 AGTGCTGTATTAGTATACTTGTAAACAGGCCACAGCCTCACCTATGACCC 318
JOURNAL	Unpublished			QY 421 TATGTAAAACTACTCCCGCCATACCAAAACCTCTTACCCACTCCGGTAC 480
REFERENCE	2 (bases 1 to 1767)			Db 317 TATGTAAAACTACTCCCGCCATACCAAAACCTCTTACCCACTCCGGTAC 258
AUTHORS	Cui, S.J., Li, Y., Li, J.W., Jin, H. and Tong, G.Z.			QY 481 TTTACCCCCAACCTGTAGATTGTTACCTTCCAAACAAACAAAGAA 540
JOURNAL	Direct Submission			Db 257 TTTACCCCCAACCTGTAGATTGTTACCTTCCAAACAAACAAAGAA 198
FEATURES	Submitted (14-JUN-2002) Harbin Veterinary Research Institute of CAAS, No. 427, Muduanjie, Nangang District, Harbin, Heilongjiang 150001, P. R. China			QY 541 ACCAGCTGTGGCTGAGACTACAAACTGCTGGAAATGTAGACCAACTGGCTGGGACT 600
Source	Location/Qualifiers			Db 197 AACAGCTGTGGCTGAGACTACAAACTGGCTGGGACT 138
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gene	/isolate="kaozha"			Db 137 GCGTTGAAAACAGTATAACGACCCAGGAATACTCGTGGCTGGGACT 78
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LOCUS	AF201897/c			
	1767 bp	DNA	circular VRL 20-DEC-2000	
	AF201897			

DEFINITION Porcine circovirus type 2, complete genome.

ACCESSION AF201897

ACCESSION AF201897.1 GI:11907587

KEYWORDS porcine circovirus type 2.

SOURCE porcine circovirus type 2. Viruses; ssDNA viruses; Circoviridae; Circovirus.

ORGANISM porcine circovirus type 2.

REFERENCE 1 (bases 1 to 1767) Wellenberg,G.J., Pesch,S., Berndsen,F.W., Steverink,P.J.G.M., Hunneman,W., Van der Vorst,T.J.K., Peperkamp,N.H.M.T., Ohlinger,V.F., Schippers,R., Van Oirschot,J.T. and de Jong,M.F.

AUTHORS

TITLE Isolation and characterization of porcine circovirus type 2 from pigs showing signs of post-weaning multisystemic wasting syndrome in The Netherlands

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 1767) Pesch,S. and Ohlinger,V.F.

AUTHORS

TITLE Direct Submission

JOURNAL Submitted (04-NOV-1999) Virology and Molecular Epidemiology, BioScreen European Veterinary Disease Management Center GmbH, 11 Mendelstr., Muenster, NRW 48149, Germany

FEATURES Location/Qualifiers

1. 1767

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51. 995

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RESULT 7  
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 LOCUS  
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 Accession AJ293869  
 Version AJ293869.1  
 Keywords  
 SOURCE  
 Organism  
 Reference  
 Authors

TITLE Isolation and characterization of porcine circovirus 2 from cases of sow abortion and porcine dermatitis and nephropathy syndrome  
 JOURNAL Arch. Virol. 146 (4), 835-842 (2001)  
 MEDLINE 21296605  
 PUBMED 11402869  
 REFERENCE 2 (bases 1 to 1766)  
 AUTHORS Meehan, B.M., McNeilly, F., McNair, I., Walker, I., West, K., Ellis, J.A., Krakowka, S., Kennedy, S., and Allan, G.M.  
 TITLE Isolation and characterization of porcine circovirus type 2 from cases of sow abortion and porcine dermatitis and nephropathy syndrome  
 JOURNAL Unpublished  
 REFERENCE 3 (bases 1 to 1766)  
 AUTHORS Meehan, B.M.  
 TITLE Direct Submission  
 JOURNAL Submitted (25-AUG-2000) Meehan B.M., Veterinary Science, The Queen's University of Belfast, Stormont, BT4 3SD, UNITED KINGDOM  
 FEATURES Location/Qualifiers  
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Query Match 97.3%; Score 682.8; DB 14; Length 1766;  
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QY 181 ACAGTCAGAACGCCCTCCTGGCCGACATGAGATTCAATTTAATGACTTTCTT 240  
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QY 1766 bp DNA circular VRL 06-JUN-2001  
 Db Porcine circovirus rep gene for replication-associated protein.  
 QY Isolate Imp.1147.  
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 QY Porcine circovirus.  
 Db Viruses; ssDNA Viruses; Circoviridae; Circovirus.  
 QY 1 (bases 1 to 1766)  
 Db Meehan, B.M., McNeilly, F., McNair, I., Walker, I., Ellis, J.A., Krakowka, S., and Allan, G.M.  
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RESULT 8  
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 KEYWORDS  
 SOURCE porcine circovirus type 2.  
 ORGANISM porcine circovirus type 2.  
 Viruses; ssDNA viruses; Circoviridae; Circovirus.  
 REFERENCE 1 (bases 1 to 1767)  
 AUTHORS Shengbo, C. and Huanchun, C.  
 TITLE Cloning and sequence analysis of the genome of Porcine Circovirus type 2 isolated from pig with PMWS in China  
 JOURNAL Unpublished  
 REFERENCE 2 (bases 1 to 1767)  
 AUTHORS Direct Submission  
 TITLE Submitted (16-MAY-2001) Animal Medicine, Hua Zhong Agricultural University, Shi Zi Shan, Wuhan, Hubei 430070, P. R. China  
 JOURNAL  
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REFERENCE 2 (bases 1 to 1768)  
 AUTHORS Hamel, A.L. and Nayar, G.P.S.  
 TITLE Nucleotide sequence of four different isolates of circovirus  
 detected in pigs with various clinical syndromes  
 unpublished  
 JOURNAL  
 REFERENCE 3 (bases 1 to 1768)  
 AUTHORS Hamel, A.L. and Nayar, G.P.S.  
 TITLE Direct Submission  
 JOURNAL Submitted (27-NOV-1998) Virology Laboratory, Veterinary Diagnostic  
 Laboratory, Manitoba Agriculture, 545 University Crescent,  
 Winnipeg, Manitoba R3T 5S6, Canada  
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 Best Local Similarity 93.0%; Pred. No. 5.6e-168;  
 Matches 653; Conservative 0; Mismatches 49; Indels 0; Gaps 0;

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Qy	481	T <sup>T</sup> TACCCCAAACCTGTCCCTAGATTCTCACTATTGATTACTTCCAACCAACAAAGA	540	Qy	181	ACAGTCACAAACGCCCTCCTGGGACTGATTGAGATTCAATAATTGACTTTCT	240
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ORGANISM				Db	1195	AATCAGCTTGGCTGAGCTAACAAACCTGGCTTACATTGATTACTTCCAACAAACAAAGA	1136
				Qy	601	GGGTTCGAAAACAGTATAACGACAGGATAACAATACCGTGTAACTGTACAA	660
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AUTHORS	KUO, T.Y., CHIOU, Y.C. and LAI, S.S.			Qy	661	TTCAGAGATTAAATTAAAGACCCCCACTTAACCCCTAA	702
TITLE	Complete nucleotide sequences analysis of porcine circovirus			Db	1075	TTCAGAGATTAAATACGACAGGACTACAATACCGTGTAACTGTACAA	1034
JOURNAL	outbreak in Taiwan						
REFERENCE	Unpublished (bases 1 to 1768)						
AUTHORS	KUO, T.Y., CHIOU, Y.C. and LAI, S.S.						
TITLE	Direct Submission						
JOURNAL	Submitted (27-MAY-1999) Veterinary Medicine, National Taiwan University, 142, Chousan Road, Taipei 106, Taiwan						
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	VGEEGNEEGRTPHLOGFANFVKQKOTENFKVWKYFARCHIEKAKGTDQDNKEYCSKEGN			TITLE	Complete nucleotide sequences of porcine circovirus Tainan strand		
	LLIECGAPRSQGQRSDLSTAVSTLLESGLSVTVAEOFHPVTFVRNFRGLAEILKVSGKM			JOURNAL	Outbreak in Taiwan		
	QKRDWKTNVHVYGGPGCKSKRWAANFADPETTYWKPRNWKWDGYHGEVVVIDFY			REFERENCE	Unpublished		
	GWLPWLDRLLCDRYPLTVEVKGGTVPFLARSILITSNQTPLEWYSSTAVPAVEALYR			AUTHORS	2 (bases 1 to 1768)		
	RITSLWVKNATEQSTEEGQFVTLSSPPCEPFYIIN"			TITLE	Yang, K.H., Lee, Y.F., Chao, D.S., Shieh, Y.C. and Lai, S.S.		
BASE COUNT	451	a	367	JOURNAL	Submitted (08-JUL-1999) Veterinary Medicine, National Taiwan		
ORIGIN			367	source	Institute of Technology, 300 Shei Fu Road, Lu Liao Li, Chia-Yi City		
			c	495	600, Taiwan		
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Best	Local	93.0%	Pred.	No. 5.6e-168;			
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ORIGIN Query Match 88.8%; Score 623.6; DB 14; Length 1768;

Best Local Similarity 93.0%; Pred. No. 5.6e-168; Matches 653; Conservative 0; Mismatches 49; Indels 0; Gaps 0;

FEATURES Source

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498 CCCCGGGAGGGGGACCAACAAATCTCAATACCCCTTGAATAAGAAAG 439

AUTHORS Mankertz, A., Domingo, M., Folch, J.M., LeCann, P., Jestin, A.,  
Segales, J., Chmielewicz, B., Plana-Duran, J. and Soike, D.

TITLE Characterisation of PCV-2 isolates from Spain, Germany and France

JOURNAL Virus Res.

66 (1), 65-77 (2000)

MEDLINE 20120936

PUBMED 10653918

REFERENCE 2 (bases 1 to 1768)

AUTHORS Mankertz, A., Domingo, M., Folch, J.M., LeCann, P., Jestin, A.,  
Segales, J., Chmielewicz, B., Plana-Duran, J. and Soike, D.

TITLE Direct Submission

JOURNAL Submitted (03-NOV-1999) P24, Robert Koch Institut, Nordufer 20,

Berlin 13353, Germany

FEATURES Location/Qualifiers

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GenCore version 5.1.6  
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Run on: May 18, 2003, 09:15:38 ; Search time 2022 Seconds

(without alignments)  
 10103.941 Million cell updates/sec

Title: US-09-514-245B-25

Perfect score: 702

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Scoring table: IDENTITY\_NUC  
 Gapop 10.0 , Gapext 1.0

Searched: 2054640 seqs, 14551402878 residues

Total number of hits satisfying chosen parameters: 4109280

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0.8

Maximum Match 100%

Listing first 45 summaries

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 41: em\_htg\_o\_other:\*

Pred. No. is the number of results predicted by chance to have a

score greater than or equal to the score of the result being printed,  
 and is derived by analysis of the total score distribution.

## SUMMARIES

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2	702	100.0	1767	6	AX003274		AX003274 Sequence
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5	698.8	99.5	1767	14	AY122275		AY122275 Porcine C
6	690.8	98.4	1767	14	AF201897		AF201897 Porcine C
7	682.8	97.3	1766	14	PC1293869		AJ293869 Porcine C
8	643.4	91.7	1767	14	AY035820		AY035820 Porcine C
9	626.8	89.3	1768	14	AF201309		AF201309 Porcine C
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11	623.6	88.8	1768	14	AF109398		AF109398 Porcine C
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27	618.8	88.1	1768	14	AF381175		AF381175 Porcine C
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35	615.6	87.7	1768	14	AF264040		AF264040 Porcine C
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37	614	87.5	1768	6	AX379562		AX379562 Sequence
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## ALIGNMENTS

RESULT 1	AX003277	LOCUS	DEFINITION	ACCESSION	VERSION	KEYWORDS	SOURCE	ORGANISM	REFERENCE
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				GI:9927101					

Porcine circovirus  
 Porcine circovirus  
 Viruses; ssDNA viruses; Circoviridae; Circovirus.  
 1 (bases 1 to 702)  
 Huet, E. and Albina, E.  
 Circovirus sequences related to piglet weight loss disease (pwd)  
 Patent: WO 9929871-A 12 17-JUN-1999;  
 HUET EVELYNE (FR); ALBINA EMMANUEL (FR)

FEATURES	Location/Qualifiers
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Db	Matches 702;保守型 0; Mismatches 0; In
Qy	1 ATGACGTATCCAAAGGAGGGCGTTACCGAAGAAAGAACCCGCC
Db	1 ATGACGTATCCAAAGGAGGGCGTTACCGAAGAAAGAACCCGCC
Qy	61 CAGATCCTCGCCGCCGCCCTGGCTCCACCCCCGCCAC
Db	61 CAGATCCTCGCCGCCGCCCTGGCTCCACCCCCGCCAC
Qy	121 AAAAATGGCATCTCAACACCCGGCTCTCCGACCTTGGAT
Db	121 AAAAATGGCATCTCAACACCCGGCTCTCCGACCTTGGAT
Qy	181 ACAGTCAGAACGCCCTGGGGACATGATGAGATTCA
Db	181 ACAGTCAGAACGCCCTGGGGACATGATGAGATTCA
Qy	241 CCCCCAGGGGGTCAAACCCCCGGCTCTGTGCCCTTGAAAT
Db	241 CCCCCAGGGGGTCAAACCCCCGGCTCTGTGCCCTTGAAAT
Qy	301 GTTAAGGTTGAATTCTGGCCCTGCTCCCCGATCACCCAGGGTG
Db	301 GTTAAGGTTGAATTCTGGCCCTGCTCCCCGATCACCCAGGGTG
Qy	361 AGTGCTGTTATTAGATAACTTTGTTAACAAAGGCCACAG
Db	361 AGTGCTGTTATTAGATAACTTTGTTAACAAAGGCCACAG
Qy	421 TATGTAAAACTACTCCCTCCGCCATACCATAACCCAGCCCTCT
Db	421 TATGTAAAACTACTCCCTCCGCCATACCATAACCCAGCCCTCT
Qy	481 TTTACCCCCAACCTGTCTAGATTTCACATTGATTACTTCC
Db	481 TTTACCCCCAACCTGTCTAGATTTCACATTGATTACTTCC
Qy	541 AACCGAGCTGTGGCTGAGACTACAAACTGCTGGAAATGTAGACC
Db	541 AACCGAGCTGTGGCTGAGACTACAAACTGCTGGAAATGTAGACC
Qy	601 GCGTTCGAAACAGTATAACGACCCAGGAATAACAAATATCCGTG
Db	601 GCGTTCGAAACAGTATAACGACCCAGGAATAACAAATATCCGTG
Qy	661 TTCAAGAGAATTAAATTAAAGACCCCCCACTTAACCCTTAA
Db	661 TTCAAGAGAATTAAATTAAAGACCCCCCACTTAACCCTTAA
RESULT 2	
AX003274 /C	
LOCUS	AX003274 1767 bp DNA
DEFINITION	Sequence 9 from Patent WO9929871.
ACCESSION	AX003274
VERSION	AX003274.1 GI:9927098
KEYWORDS	porcine circovirus. porcine circovirus Viruses; ssDNA viruses; Circoviridae; Circovir
SOURCE	1 (bases 1 to 1767) Hutet, E. and Albina, E.
ORGANISM	
REFERENCE	
AUTHORS	

KW Porcine multisystemic wasting syndrome; pig; vaccine; ss.  
 XX Porcine circovirus.  
 XX PN FR2769322-A1.  
 XX PD 09-APR-1999.  
 XX PF 22-JAN-1998; 98FR-0000873.  
 XX PR 03-OCT-1997; 97FR-0012382.  
 XX PA (MERI-) MERIAL SAS.

XX PI Allan G, Chappuis GE, Charreyre CE, Clark E, Ellis J;  
 PI Haines D, Harding J, Hassard L, Meehan B;  
 XX DR WPI: 1999-246948/21.  
 XX PT New type II porcine circovirus, used for, e.g. passive immunization  
 PT of pregnant sows  
 XX PS Claim 14; Fig 2; 48pp; French.

XX The present sequence represents the nucleotide sequence of PCV isolate  
 CC Imp1011-48285. The specification describes a preparation of type II  
 CC porcine circovirus (PCV), which is particularly isolated from a lesion,  
 CC from a pig with symptoms of PMWS (porcine multisystemic wasting  
 CC syndrome). PCV (attenuated or inactivated), polypeptides derived from  
 CC it, and vectors that express these polypeptides are all useful in  
 CC vaccines, suitable for administration to adult or young pigs, or to  
 CC pregnant sows (for passive immunization of their offspring). DNA  
 CC isolated from PCV is used for in vivo or in vitro expression of viral  
 CC polypeptides, also as probes or primers for diagnosis in usual  
 CC hybridization or amplification assays. These polypeptides may also be  
 CC used diagnostically to detect PCV-specific antibodies, while antibodies  
 CC raised against the polypeptides can be used to detect antigens, in any  
 CC usual immunoassay format.

XX SQ Sequence 1767 BP; 448 A; 359 C; 500 G; 460 T; 0 other.

Query	Match	Score	Length						
Best	Local	Similarity	Pred.	No.	Mismatches	Indels	Gaps	0;	
QY	315	CTGGCCCTGCTCCCCGATCACCCAGGGTACAGGGGAGTGGCTCCAGTGCTGTATT	53.9%	378.4	DB 20;	Length 1767;			
Db	1767	CTGGCCCTGCTCCCCGATCACCCAGGGTACAGGGGAGTGGCTCCAGTGCTGTATTCT	98.5%	5.2e-104;					
QY	375	AGATGATAACTTTGTAACAAAGGCCACAGGCCCTCACCTATGACCCCTATGTAAACTACTC	0;						
Db	1707	AGATGATAACTTTGTAACAAAGGCCACAGGCCCTCACCTATGACCCCTATGTAAACTACTC	0;						
QY	435	CTCCCGCCATACCATAACCCAGCCCTTCTCTACCATCTCCCGTACTTTACCCCCAAC	374	4.34					
Db	1647	CTCCCGCCATACCATAACCCAGCCCTTCTCTACCATCTCCCGTACTTTACCCCCAAC	1588	1.648					
QY	495	TGTCTAGATTTCACCTATTGATTACTTCCGTTAACCATGTACAATTGAGAAATTAA	494	4.94					
Db	1587	TGTCTAGATTCCACATTGATTACTTCCAAACCAACAAACAAATCAGCTGTGGCT	554	1.528					
QY	555	GAGACTACAAACTGCTGAAATGTAGACCCACGTAGGGCTCGGGCACTGCGTT	614	6.14					
Db	1527	GAGACTACAAACTGTTAACCATGTACAATTGAGCCACGTAGGGCTCGGGCACTGCGTT	1468	1.468					
QY	615	TATATAGACCAGGAATACAATATCCGTGTAACCATGTACAATTGAGAAATTAA	674	6.74					
Db	1467	TATATACGGACCAGGAATACAATATCCGTGTAACCATGTACAATTGAGAAATTAA	1408	1.408					
QY	675	TTTTAAGACCCCCCACTTAACCCCTAA	702	7.02					
Db	1407	TCTTAAGACCCCCCACTTAACCCCTAA	1380	1.380					

Best Local Similarity 99.0%; Pred. No. 5.6e-105;	Indels 0;	Gaps 0;	CC isolate.
Matches 384; Conservative 0; Mismatches 4;	Sequence 1767 BP; 447 A; 360 C; 502 G; 458 T; 0 other;	XX	XX
QY 315 CTGGCCCTGCTCCCGATCACCCAGGGTACAGGGGAGTGGCTCCAGTGGCTATT 374	Query Match 54.4%; Score 381.6; Pred. No. 5.6e-105;	XX	XX
Db 1767 CTGGCCCTGCTCCCGATCACCCAGGGTACAGGGGAGTGGCTCCAGTGGCTATT 1708	Best Local Similarity 99.0%; Mismatches 0; Indels 0; Gaps 0;	XX	XX
QY 375 AGATGATAACTTTGTAACAAAGGCCAACAGCCCTCACCTATGACCCCTATGTAACACTC 434	Matches 384; Conservative 0; Mismatches 4;	XX	XX
Db 1707 AGATGATAACTTTGTAACAAAGGCCAACAGCCCTCACCTATGACCCCTATGTAACACTC 1648	Query 315 CTGGCCCTGCTCCCGATCACCCAGGGTACAGGGGAGTGGCTCCAGTGGCTATT 374	XX	XX
QY 435 CTCGGCCATACCAAAACCCAGCCCTTCTCCTACCACTCCGGTACTTTACCCCCAAC 94	Db 1767 CTGGCCCTGCTCCCGATCACCCAGGGTACAGGGGAGTGGCTCCAGTGGCTATT 1708	XX	XX
Db 1647 CTCGGCCATACCAAAACCCAGCCCTTCTCCTACCACTCCGGTACTTTACCCCCAAC 1588	QY 375 AGATGATAACTTTGTAACAAAGGCCAACAGCCCTCACCTATGACCCCTATGTAACACTC 434	XX	XX
QY 495 TGTCTTAGATTTCACCTATTGATTACTTCCAAACAAACAAACAAAGAACAGCTGGCT 554	Db 1707 AGATGATAACTTTGTAACAAAGGCCAACAGCCCTCACCTATGACCCCTATGTAACACTC 1648	XX	XX
Db 1587 TGTCTTAGATTTCACCTATTGATTACTTCCAAACCAACAAACAAAGAACAGCTGGCT 1528	QY 435 CTGGCCCTGCTCCCGATCACCTTCTCCTACCACTCCGGTACTTTACCCCCAAC 94	XX	XX
QY 555 GAGACTACAAACTGCTGGAAATGTAGACCCACGTAGGGCCTCGGCACTGGTTCGAAAACAG 614	Db 1647 CTGGCCCTGCTCCCGATCACCTTCTCCTACCACTCCGGTACTTTACCCCCAAC 1588	XX	XX
Db 1527 GAGACTACAAACTGCTGGAAATGTAGACCCACGTAGGGCCTCGGCACTGGTTCGAAAACAG 1468	QY 495 TGTCTTAGATTTCACCTATTGATTACTTCCAAACAAACAAACAAAGAACAGCTGGCT 554	XX	XX
QY 615 TATATACGACCAATAACAAATACTCGTAGACCATGTTATGTACAAATTTCAGAGAATTAA 674	Db 1587 TGTCTTAGATTTCACCTATTGATTACTTCCAAACAAACAAACAAAGAACAGCTGGCT 1528	XX	XX
Db 1467 TATATACGACCAATAACAAATACTCGTAGACCCACGTAGGGCTACCAATTTCAGAGAATTAA 1408	QY 555 GAGACTACAAACTGCTGGAAATGTAGACCCACGTAGGGCCTCGGCACTGGCTCGAAAACAG 614	XX	XX
QY 675 TTTAAAGACCCCCCACTTAACCCCTAA 702	Db 1527 GAGACTACAAACTGCTGGAAATGTAGACCCACGTAGGGCCTCGGCACTGGCTCGAAAACAG 1468	XX	XX
Db 1407 TCTTAAGACCCCCCACTTAACCCCTAA 1380	QY 615 TATATACGACCAATAACAAATACTCGTAGACCATGTTATGTACAAATTTCAGAGAATTAA 674	XX	XX
RESULT 1.2	Db 1467 TATATACGACCAATAACAAATACTCGTAGACCATGTTATGTACAAATTTCAGAGAATTAA 1408	XX	XX
AAZ56869/C	QY 675 TTTAAAGACCCCCCACTTAACCCCTAA 702	XX	XX
ID AAZ56869 standard; DNA; 1767 BP.	Db 1407 TCTTAAGACCCCCCACTTAACCCCTAA 1380	XX	XX
XX	RESULT 1.3	XX	XX
AC AAZ56869;	AAF75835/C	XX	XX
XX	ID AAF75835 standard; DNA; 1767 BP.	AC	AC
DT 25-APR-2000 (first entry)	XX	XX	XX
DE DNA sequence of PCV Imp.1011-48121 isolate.	XX	XX	XX
XX	XX	XX	XX
DE. DNA sequence of PCV Imp.1011-48121 isolate.	XX	XX	XX
XX	KW Antigen; porcine multisystemic wasting syndrome; PMWS; antiviral;	XX	XX
KW porcine circovirus; PCV; porcine parvovirus; PPV; vaccination; ds.	XX	XX	XX
XX	OS Porcine circovirus.	XX	XX
OS	XX	XX	XX
PN WO200001409-A2.	XX	XX	XX
XX	OS Porcine circovirus-2.	XX	XX
PD 13-JAN-2000.	XX	XX	XX
XX	OS Porcine circovirus-2.	XX	XX
PF 28-JUN-1999; 99WO-EP04698.	XX	XX	XX
XX	PF 28-AUG-2000; 2000WO-EP08781.	XX	XX
PR 06-JUL-1998; 98FR-0008777.	XX	XX	XX
XX	PR 31-AUG-1999; 99US-0151564.	XX	XX
PA (MERI-) MERIAL.	XX	XX	XX
PA (UYBE-) UNIV QUEENS BELFAST.	XX	XX	XX
PA (UYSA-) UNIV SASKATCHEWAN.	XX	XX	XX
XX	PR 31-MAY-2000; 2000US-0583350.	XX	XX
PI Allan GM, Meehan BM, Ellis JA, Krakowka GS, Audonnet JF;	PA (MERI-) MERIAL.	XX	XX
XX	PA (UYSA-) UNIV SASKATCHEWAN.	XX	XX
DR WPI; 2000-182091/16.	PA (UYBE-) UNIV QUEENS BELFAST.	XX	XX
XX	PI Harding J, Charreyre CE, Chappuis GE, Krakowka GS, Haines D, Audonnet JF;	XX	XX
PT Use of a porcine circovirus antigen and a porcine parvovirus antigen for vaccination against porcine multisystemic wasting syndrome -	PI Mcneilly F;	XX	XX
PT	XX	XX	XX
PS Disclosure; Fig 1; 61PP; English.	XX	XX	XX
XX	XX	XX	XX
CC The invention provides a novel antigenic preparation directed against porcine multisystemic wasting syndrome (PMWS) that comprises porcine circovirus (PCV) antigen and porcine parvovirus (PPV) antigen. The PCV antigens and PPV antigens can be used for vaccination against PMWS. The present sequence represents the DNA sequence of PCV Imp.1011-48121	DR	XX	XX
CC	XX	XX	XX
CC	XX	XX	XX
CC	XX	XX	XX
CC	XX	XX	XX

PT infection and/or post-weaning, multisystemic wasting syndrome  
 PT associated with PCV-2  
 XX Disclosure; Fig. 1 #2; 134pp; English.

CC The present invention relates to the use of porcine circovirus-2 (PCV-2)  
 CC immunogen to formulate a vaccine composition to prevent or treat pigs  
 CC against myocarditis and/or abortion and/or intrauterine infection and/or  
 CC post-weaning, multisystemic wasting syndrome and other pathological  
 CC sequelae associated with PCV-2. The present sequence is a DNA fragment of  
 CC a strain of PCV, which was used in the present invention.

XX Sequence 1767 BP; 447 A; 360 C; 502 G; 458 T; 0 other;

Query Match 54.4%; Score 381.6; DB 22; Length 1767;  
 Best Local Similarity 99.0%; Pred. No. 5.6e-105;  
 Matches 384; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

QY 315 CTGGCCCTGCTCCCGATCACCCAGGGTACAGGGGACTGGGTCCAGTGGCTTATT 374  
 Db 1767 CTGGCCCTGCTCCCGATCACCCAGGGTACAGGGGACTGGGTCCAGTGGCTTATTCT 1708

QY 375 AGATGATAACTTTGTAAACAAAGGGCCACAGGCCCTACCTATGACCCCTATGTAACACTC 434  
 Db 1707 AGATGATAACTTTGTAAACAAAGGGCCACAGGCCCTACCTATGACCCCTATGTAACACTC 1648

OY 435 CTCGGCCATACCATAACCCAGCCCTTCTCCTACCACTCCCCGTACTTTACCCCCAACCC 494  
 Db 1647 CTCCGGCCATACCATAACCCAGCCCTTCTCCTACCACTCCCCGTACTTTACCCCCAACCC 1588

QY 495 TGTCCCTAGATTTCACATTGATTACTTCCAACCAAACAAAGAAACCCAGCTGTGGCT 554  
 Db 1587 TGTCCCTAGATTTCACATTGATTACTTCCAACCAAACAAAGAAACCCAGCTGTGGCT 1528

QY 555 GAGACTACAAACTGCTGGAAATGTAGACCACTGGCCTAGGTTCGAAACAG 614  
 Db 1527 GAGACTACAAACTGCTGGAAATGTAGACCACTGGCCTAGGTTCGAAACAG 1468

QY 615 TATATAGGACCGAAATACAATATCCGTTAACCATGTTACCAATTCAAGAGAATTAA 674  
 Db 1467 TATATAGGACCGAAATACAATATCCGTTAACCATGTTACCAATTCAAGAGAATTAA 1408

QY 675 TTTAAAGACCCCCCACTTAACCCCTTA 702  
 Db 1407 TCTTAAAGACCCCCCACTTAACCCCTTA 1380

RESULT 14  
 ID AAX35379/C  
 XX 07-JUL-1999 (first entry)  
 DE Nucleotide sequence of PCV isolate Imp1011-48285.  
 XX PCV isolate; type II porcine circovirus; PCV; PMWS;  
 KW porcine multisystemic wasting syndrome; pig; vaccine; ss.  
 XX Porcine circovirus.  
 XX PN W09918214-A1.  
 XX PD 15-APR-1999.  
 XX PF 01-OCT-1998; 98WO-FR02107.  
 XX PR 20-MAR-1998; 98FR-0003707.  
 PR 03-OCT-1997; 97FR-0012382.  
 PR 22-JAN-1998; 98FR-000873.  
 PA (MERI-) Merial.

PA (UYBE-) UNIV QUEENS BELFAST.  
 PA (UYSA-) UNIV SASKATCHEWAN.  
 XX PI Allan G, Chappuis GE, Charreyre CE, Clark E, Ellis J;  
 PI Haines D, Harding J, Hassard L, Meehan B;  
 XX DR WPI; 1999-264024/22.  
 PT New type II porcine circovirus  
 XX PS Claim 11; Fig 2; 56pp; French.  
 XX  
 The present sequence represents the nucleotide sequence of PCV isolate  
 CC Imp1011-48285. The specification describes a preparation of type II  
 CC porcine circovirus (PCV), which is particularly isolated from a lesion,  
 CC from a pig with symptoms of PMWS (porcine multisystemic wasting  
 syndrome). PCV (attenuated or inactivated), polypeptides derived from  
 CC it, and vectors that express these polypeptides are all useful in  
 CC vaccines, suitable for administration to adult or young pigs, or to  
 CC pregnant sows (for passive immunization of their offspring). DNA  
 CC isolated from PCV is used for in vivo or in vitro expression of viral  
 CC polypeptides, also as probes or primers for diagnosis in usual  
 CC hybridization or amplification assays. These polypeptides may also be  
 CC used diagnostically to detect PCV-specific antibodies, while antibodies  
 CC raised against the polypeptides can be used to detect antigens, in any  
 CC usual immunoassay format.  
 XX Sequence 1767 BP; 448 A; 359 C; 500 G; 460 T; 0 other;  
 SQ Query Match 53.9%; Score 378.4; DB 20; Length 1767;  
 Best Local Similarity 98.5%; Pred. No. 5.2e-104;  
 Matches 382; Conservative 0; Mismatches 6; Indels 0; Gaps 0;

QY 315 CTGGCCCTGCTCCCGATCACCCAGGGTACCCAGGGTACAGGGGACTGGGTCCAGTGGCTTATT 374  
 Db 1767 CTGGCCCTGCTCCCGATCACCCAGGGTACCCAGGGTACAGGGGACTGGGTCCAGTGGCTTATTCT 1708

QY 375 AGATGATAACTTTGTAAACAAAGGGCCACAGGCCCTACCTATGACCCCTATGTAACACTC 434  
 Db 1707 AGATGATAACTTTGTAAACAAAGGGCCACAGGCCCTACCTATGACCCCTATGTAACACTC 1648

QY 435 CTCCGGCCATACCATAACCCAGCCCTTCTCCTACCACTCCCCGTACTTTACCCCCAACCC 494  
 Db 1647 CTCCGGCCATACCATAACCCAGCCCTTCTCCTACCACTCCCCGTACTTTACCCCCAACCC 1588

QY 495 TGTCCCTAGATTTCACATTGATTACTTCCAACCAAACAAAGAAACCCAGCTGTGGCT 554  
 Db 1587 TGTCCCTAGATTTCACATTGATTACTTCCAACCAAACAAAGAAACCCAGCTGTGGCT 1528

QY 555 GAGACTACAAACTGCTGGAAATGTAGACCACTGGCCTAGGTTCGAAACAG 614  
 Db 1527 GAGACTACAAACTGCTGGAAATGTAGACCACTGGCCTAGGTTCGAAACAG 1468

QY 615 TATATAGGACCGAAATACAATATCCGTTAACCATGTTACCAATTCAAGAGAATTAA 674  
 Db 1467 TATATAGGACCGAAATACAATATCCGTTAACCATGTTACCAATTCAAGAGAATTAA 1408

QY 675 TTTAAAGACCCCCCACTTAACCCCTTA 702  
 Db 1407 TCTTAAAGACCCCCCACTTAACCCCTTA 1380

RESULT 15  
 ID AAX35211/C  
 XX 01-JUL-1999 (first entry)  
 DE Nucleotide sequence of PCV isolate Imp1011-48285.  
 XX PCV isolate; type II porcine circovirus; PCV; PMWS;  
 KW porcine multisystemic wasting syndrome; pig; vaccine; ss.  
 XX Porcine circovirus.  
 XX PN W09918214-A1.  
 XX ID AAX35211 standard; DNA; 1767 BP.  
 XX AC AAX35211;  
 XX DT 01-JUL-1999 (first entry)  
 XX DE Nucleotide sequence of PCV isolate Imp1011-48285.  
 XX KW PCV isolate; type II porcine circovirus; PCV; PMWS;

QY	653	ATGTAACATTAGAGAATTAAATTAAAGACCCCACTTAACCCCTTAA	702	Db	1647	CTCCGCCATACCACTGGCCttCTCTACTCCGGTACTttACCCCAAAAC	1588		
Db	664	ATGTAACATTAGAGAATTAAATTAAAGACCCCACTTAACCCCTAA	615	QY	495	TGTCCTAGATTCACTTATTGATTAACCAACAAAGAAACCAGtGTGGCT	554		
<b>RESULT 10</b>									
ID	AAX35378/C	standard; DNA; 1767 BP.		Db	1587	TGTCCTAGATTCCACTTATTGATTAACCAACAAACAAAGAAACCAGtGTGGCT	1528		
XX	AAX35378;			QY	555	GAGACTACAAACTGCTGGAAATGTAGACCACTGGCACTGGTTCGAAAACAG	614		
XX	DT 07-JUL-1999	(first entry)		Db	1527	GAGACTACAAACTGCTGGAAATGTAGACCACTGGCACTGGTTCGAAAACAG	1468		
XX	DE	Nucleotide sequence of PCV isolate Imp1011-48121.		QY	615	TATATACGACCAAGGAAATACAATATCCGTTAACCATGTTACAAATTCAAGGAATTAA	674		
XX	KW	PCV isolate; type II porcine circovirus; PCV; PMWS;		Db	1467	TATATACGACCAAGGAAATACAATATCCGTTAACCATGTTACAAATTCAAGGAATTAA	1408		
XX	KW	porcine multisystemic wasting syndrome; pig; vaccine; ss.		QY	675	TTTTAAAGACCCCCACTTAACCCCTAA	702		
XX	OS	porcine circovirus.		Db	1407	TCTTAAGACCCCCACTTAACCCCTAA	1380		
XX	PN	W09918214-A1.		<b>RESULT 11</b>					
XX	XX	AAX35210/C		ID	AAX35210	standard; DNA; 1767 BP.			
PD	15-APR-1999.			XX	AAX35210;				
XX	PF 01-OCT-1998;	98WO-FR02107.		AC					
XX	PR 20-MAR-1998;	98FR-0003707.		XX		01-JUL-1999 (first entry)			
PR	03-OCT-1997;	97FR-0012382.		DT					
PR	22-JAN-1998;	98FR-0000873.		XX					
XX	PA (MERI-) MERIAL.			DE					
PA (UYBE-) UNIV QUEENS BELFAST.				XX					
PA (UYSA-) UNIV SASKATCHEWAN.				KW					
XX	PI Allan G, Chappuis GE, Charreyre CE, Clark E, Ellis J;			KW					
PI Haines D, Harding J, Hassard L, Meehan B;				XX					
XX	DR WPI: 1999-264024/22.			OS					
XX	PT New type II porcine circovirus			OS					
XX	PS Claim 11; Fig 1; 56pp; French.			XX					
XX	CC The present sequence represents the nucleotide sequence of PCV isolate			XX					
CC Imp1011-48121. The specification describes a preparation of type II				PA					
CC porcine circovirus (PCV), which is particularly isolated from a lesion,				XX					
CC from a pig with symptoms of PMWS (porcine multisystemic wasting				PI					
CC syndrome). PCV (attenuated or inactivated), polypeptides derived from				PI					
CC it, and vectors that express these polypeptides are all useful in				Haines D, Chappuis GE, Charreyre CE, Clark E, Ellis J;					
CC vaccines, suitable for administration to adult or young pigs, or to				XX					
CC pregnant sows (for passive immunization of their offspring). DNA				DR					
CC isolated from PCV is used for in vivo or in vitro expression of viral				WPI: 1999-246948/21.					
CC polypeptides, also as probes or primers for diagnosis in usual				XX					
CC hybridization or amplification assays. These polypeptides may also be				XX					
CC used diagnostically to detect PCV-specific antibodies, while antibodies				PA					
CC raised against the polypeptides can be used to detect antigens, in any				XX					
CC usual immunoassay format.				CC					
XX	Sequence 1767 BP; 447 A; 360 C; 502 G; 458 T; 0 other;			CC					
SQ	Query Match 54.4%; Score 381.6; DB 20; Length 1767;			CC					
	Best Local Similarity 99.0%; Pred. No. 5.6e-105;			CC					
Matches 384; Conservative 0; Mismatches 4; Indels 0; Gaps 0;				CC					
QY	315 CTGGCCCTGCTCCCGATCACCCAGGGTACAGGGGAGTGGGCTCCAGTGGCTTATT	374		CC					
Db	1767 CTGGCCCTGCTCCCGATCACCCAGGGTACAGGGGAGTGGGCTCCAGTGGCTTATT	1708		CC					
QY	375 AGATGATAACTTGTAAACAAAGGCCACAGCCCTCACCTATGACCCCTATGTAAACTACTC	434		CC					
Db	1707 AGATGATAACTTGTAAACAAAGGCCACAGCCCTCACCTATGACCCCTATGTAAACTACTC	1648		CC					
QY	435 CTCCGGCATACCATAACCCAGCCCTTCTACCACTCCGGTACTTACCCCAAACC	494		XX					
	Query Match 54.4%; Score 381.6; DB 20; Length 1767;			XX					

Best Local Similarity	92.0%	Pred.	No. 8.6e-170;				
Matches	653; Conservative 0; Mismatches 49;	Indels	8;	Gaps	2;		
QY	1 ATGACGTATCCAAGGAGGGCGTACCGAAGAAGAACCCGCCGCAGCCATCTTGGC	60		PR	16-DEC-1997;	97US-0069750.	
Db	1749 ATGACGTATCCAAGGAGGGCGTACCGAAGAAGAACCCGCCGCAGCCATCTTGGC	1690		PR	11-DEC-1997;	97US-0069233.	
QY	61 CAGATCCCTGGCCGCCCTGGCTCGTCCACCCCGCC-----ACCGTTACCGCTG	113		XX	(UYSA-)	UNIV SASKATCHEWAN.	
Db	1689 CAGATCCCTGGCCGCCCTGGCTCGTCCACCCCGCCCTACCGCTACCGCTG	1630		PA			
QY	114 GAGAAGGAAAATGGCATCTTCAACACCCGCCCTCCCGCACCTCGGATAACTGTCAA	173		XX	PI	Babiuk LA, Potter AA, Wang L, Willson P;	
Db	1629 GAGAAGGAAAATGGCATCTTCAACACCCGCCCTCCCGCACCTCGGATAACTGTCAA	1570		PT	XX	New isolated porcine circovirus type II	
QY	174 GCGAACCCAGTCGAACGGCCCTCTGGGGCATGATGAGATTCAATTAAATGA	233		PS	XX	Claim 1; Fig 4; 82pp; English.	
Db	1569 GCGTACCAAGTCACAACGGCCCTCTGGGGCATGATGAGATTAAATTGACGA	1510		CC		The present invention describes a new isolated porcine circovirus type II (PCVII), obtained from postweaning multisystemic wasting syndrome-affected pigs. AAY83754 to AAY83757 represent PCVII nucleotide sequences. AAY24939 to AAY24934 represent PCVII open reading frame (ORF) proteins (N.B. the PCVII ORFs given in Fig 2A to Fig 2B do not correspond exactly with the PCVII ORFs given in Fig 3A to Fig 3D).	
QY	234 CTTTCTTCCCCAGGAGGGGTCAAACCCCCGCTCTGGCCCTTGAATACTACAGAT	293		CC	CC	The PCVII polypeptides can be used for treating or preventing PCVII infection in vertebrates. The products can also be used to detect the PCVII.	
Db	1509 CTTGTTCCCCGGAGGGGGACCAACAAAATCTCTATAACCCCTGAATACTACAGAT	1450		CC	CC		
QY	294 AAGAAAGGTTAAGGGTTGAATTCTGGCCCTGCTCCCCGATCACCCAGGGGAGT	353	Sequence	1361	BP;	334 A; 265 C; 375 G; 387 T; 0 other;	
Db	1449 AAGAAAGGTTAAGGGTTGAATTCTGGCCCTGCTCCCCGATCACCCAGGGGAGT	1390	Query	84.2%	Score	590.8; DB 20;	Length 1361;
QY	354 GGGCTCCAGTGCTGTATTAGTATAACTTTGTAAACAAAGGCCACAGCCCTCACCTA	413	Best Local Similarity	91.5%	Pred.	No. 2.1e-168;	
Db	1389 GGGCTCCACTGCTGTATTAGTATACTTTGTAAACAAAGGCCACAGCCCTAACCTA	1330	Matches	650;	Mismatches	0; Indels	2;
QY	414 TGACCCCTATGTAACACTCTCCGCCATACCATAACCCAGCCCTTCTCTACACTC	473	QY	1	ATGACGTATCCAAGGAGGGTACCGAAGAAGAACCCGCCAGCCATCTTGGC	60	
Db	1329 TGACCCATATGTAACACTCTCCGCCATACCATCCCCAACCTCTCTACACTC	1270	Db	1324	ATGACGTATCCAAGGAGGGTACCGAAGAAGAACCCGCCAGCCATCTTGGC	1265	
QY	474 CCGGTACTTTACCCCCAACCTGTCTAGATTTCACATTGATACTTCACACTC	533	QY	61	CAGATCCTCCGCCGCGCCCTGGCTCGTCACCCCCGCC-----ACCGTTACCGCTG	113	
Db	1269 CGGTTACTTCACACCCAAACCTGTCTTGACTCCACTATTGATTACTTCACAAATAA	1210	Db	1264	CAGATCCTCCGCCGCGCCCTGGCTCGTCACCCCCGCCCTACCGTTACCGCTG	1205	
QY	534 CAAAGAAACCAACCGCTGTGGCTGAGACTACAAACTGTAGACCCGTAGGGCT	593	QY	114	GAGAAGGAAAATGGCATCTTCAACACCCGCTCTCCGCACCTTTCGGATATACTGTCAA	173	
Db	1209 CAAAGGAATCAGCTTTGGCTGAGGCTACAAACCTCTGGAAATGGACCATGCT	1150	Db	1204	GAGAAGGAAAATGGCATCTTCAACACCCGCTCTCCGCACCTTTCGGATATACTGTCAA	1145	
QY	594 CGGCACTGGTTCGAAACAGTATAAC-GACCAGGAATAAACTAACCTAAACCATGT	652	QY	174	GCGAACCAAGTCAGAACGGGGCTCCTGGGGTGGACATGATGAGATTCAGAGAT	233	
Db	1149 CGGCACTGGTTCGAAACAGTATAAC-GACCAGGAATAAACTAACCTAAACCATGT	1090	Db	1144	AGCTACCAAGTCACAACGCCCTCTGGGGTGGACATGAGATTAAATTGACGA	1085	
QY	653 ATGTACAATTCAAGAATTAACTTAAAGACCCCCACTAACCTTAAACCTTAA	702	QY	234	CTTTCTCCCCAGGAGGGGTCAAACCCGGCTCTGGCCCTGCTCCCCGATCACCCCTTGAATACTACAGAA	293	
Db	1089 ATGTACAATTCAAGAATTAACTTAAAGACCCCCACTAACCTTAAACCTTAA	1040	Db	1084	CTTTGTTCCCCGGAGGGGACCAAAATCTCTATAACCCCTTGAATACTACAGAA	1025	
QY	RESULT <sup>9</sup>		QY	294	AAGAAAGGTTAAGGGTTGAATTCTGGCCCTGCTCCCCGATCACCCAGGGGAGT	353	
ID	AAX83757/c		QY	354	GGGCTCCAGGTGTATTAGATGATAACTTGTAAACAAAGGCCACAGCCCTAACCTA	413	
XX	ATX83757	standard; DNA; 1361 BP.	Db	964	GGGCTCCACTGCTGTATTAGATGATAACTTGTAAACAAAGGCCACAGCCCTAACCTA	905	
AC	AAX83757;		QY	414	TGACCCCTATGTAACACTCTCCGCCATACCATCCCCAACCCCTCTCCCTACACTC	473	
XX	DT 27-AUG-1999 (first entry)		Db	904	TGACCCATATGTAACACTCTCCGCCATACCATCCCCAACCCCTCTCCCTACACTC	845	
DE	Porcine circovirus type II B9 nucleotide sequence fragment #2.		QY	474	CCGGTACTTTACCCCCAACCTGTCTAGATTGATTACTTCACATTGATTACTTCAC	533	
XX	KW Porcine circovirus type II; PCVII; pig; infection; vaccine; postweaning multisystemic wasting syndrome virus; diagnosis; ds.		Db	844	CCGGTACTTCACACCCAAACCTGTCTTGACTCCACTATTGATTACTTCAC	785	
XX	OS Porcine circovirus.		QY	534	CAAAGAAACCAACGGCTGTGGCTGAGACTACAAACTGTGAAATGTAGACCAACTGTTAGGGCT	593	
XX	PN WO929717-A2.		Db	784	CAAAAGGAATCAGCTTTGGCTGAGGCTACAAACCTCTGGACACGTAGGGCT	725	
XX	PD 17-JUN-1999.		QY	594	CGGCACACTGGTTCGAAAACAGTATAAC-GACCAGGAATAACATCCGTGTAACCATG	652	
XX	PF 11-DEC-1998; 98WO-CA01130.		Db	724	CGGCACACTGGTTCGAAAACAGTAAATAACAGACCAGGACTACAAATCCGTGTAACCATG	665	

1615	AAAATGGCATTTCAACACCCGGCTCTCCGCACCTTCAGGATAACTGTCAAGCGTACC	1556
181	ACAGTCAGAACGCCCTCCTGGGGTGGGACATGATGAGATTCAATAATTGACTTCTT	240
1555	ACAGTCACAACGCCCTCCTGGGGTGGGACATGATGAGATTAAATTGACCGACTTGT	1496
241	CCCCCAGGGGGTCAAACCCCCGGCTCTGGCCCTTGTGAATACTACAGAATAAGAAAG	300
1495	CCCCGGGGAGGGACCAACAAATCTCTATAACCCCTTGAATACTACAGAATAAGAAAG	1436
301	GTAAAGGTTGAATTCTGGCCCTGCTCCCCGATCACCCAGGGTGAACAGGGAGTGGCTCC	360
1435	GTAAAGGTTGAATTCTGGCCTTGCTCCCCCATCACCCAGGGTGAATAGGGAGTGGCTCC	1376
361	AGTGCTGTTATTTAGATGATAACTTGTAAACAAGGCCACAGGCCCACACAGGCCACAGGCCAACAAAGGGTAACTTGTAAACAACCCATAACCTATGACCCC	420
1375	ACTGCTGTTATTTAGATGATAACTTGTAAACAAGGGTAAACAAAGGCCACAGGCCACAGGCCAACAAAGGGTAACTTGTAAACAACCCATAACCTATGACCCA	1316
421	TATGTAAACTACTCCCGCATAACCATAACCCAGCCCTTCTCCTTACCACTCCGGTAC	480
1315	TATGTAAACTACTCCCTCCGGCATAACCATCCCCAACCCCTTCTCCTTACCACTCCGGTAC	1256
481	TTTACCCCCAACCTGCTTAGATTCACTATTGATTACTTCCAACCAAACAAAGA	540
1255	TTCACACCCAACCTGTTGACTCCACTATTGATTACTTCCAACCAAATAACAAAGG	1196
541	AACCAGCTGTGGCTGAGACTACAACCTGCTGGAAATGTAGACCACGTAGGGCACT	600
1195	AATCAGCTTGGCTGAGGCTACAAACCTCTGGAAATGTGGACCACGTAGGGCACT	1136
601	GGGTTCGAAACAGTATAACGACCGGAATAACATTATCCGTGTAACCATGTACAA	660
1135	GGGTTCGAAACAGTAAATAGACCGGACTACAAATCCGTGTAACCATGTACAA	1076
661	TTCAGAGAATTAAATTAAAGACCCCCACTTAACCCCTTAA	702
1075	TTCAGAGAATTAAATCTTAAAGACCCCCACTTGAACCCCTTAA	1034

9 TESCHI

AF75841 standard; DNA; 1768 BP.

22276841

118-MAY-2001 (first entry)  
AAF/3841;

PCV DNA fragment of Imp 1121 strain.

vaccine; py; myocarditis; abortion; intrauterine infection; multisystemic wasting syndrome; ds.

Porcine circovirus-2.

WO2008116330-A2.

08 - MAR - 2001 .

28 - NYC - 30000 : 3000000 - ER08781

31-AUG-1999; 99US-0151564.  
28-APR-2000; 2000W0-E008/81.

31-MAY-2000; 20000US-0583350.

(MERIT) MERIAH

(U.S.A.) UNIV SASKATCHEWAN.

(UYBE-) UNIV QUEENS BELFAST.

卷之三

RESULT /  
AAF75840/C  
ID AAF75840 standard; DNA; 1768 BP.  
XY

AC	AAF75840;	XX	361	AGTGCTGTTATTAGATGATAACTTGTAAACAAGGCCACAGGCCCTCACCTATGACCCC	420
	18-MAY-2001	(first entry)			
DE	PCV DNA fragment of Imp 1103 strain.	XX			
XX	Vaccine; pig; myocarditis; abortion; intrauterine infection; multisystemic wasting syndrome; ds.	XX			
XX	Porcine circovirus-2.	XX			
XX	WO200116330-A2.	XX			
PD	08-MAR-2001.	XX			
XX	28-AUG-2000; 2000WO-EP08781.	XX			
XX	PR 31-AUG-1999; 99US-0151564.	XX			
XX	PR 31-MAY-2000; 2000US-0583350.	XX			
PA	(MERI-) MERIAL.	XX			
PA	(UUSA-) UNIV SASKATCHEWAN.	XX			
PA	(UYBE-) UNIV QUEENS BELFAST.	XX			
PI	Ellis JA, Allan GM, Meehan B, Clark E, Haines D, Hassard L; Harding J, Charreyre CE, Chappuis GE, Krakowka GS, Audonnet JF; Mcneilly F;	XX			
XX	WPI; 2001-244408/25.	XX			
XX	PT Use of porcine circovirus-2 immunogen to formulate a vaccine	XX			
PT	composition to treat pigs against myocarditis, abortion, intrauterine	XX			
PT	infection and/or post-weaning, multisystemic wasting syndrome	XX			
PT	associated with PCV-2.	XX			
PS	Claim 10; Fig 6 #2; 134pp; English.	XX			
XX	The present invention relates to the use of porcine circovirus-2 (PCV-2) immunogen to formulate a vaccine composition to prevent or treat pigs against myocarditis and/or abortion and/or intrauterine infection and/or post-weaning, multisystemic wasting syndrome and other pathological sequelae associated with PCV-2. The present sequence is a DNA fragment of a strain of PCV, which was used in the present invention.	XX			
SQ	Sequence 1768 BP; 450 A; 360 C; 496 G; 460 T; 2 other;	XX			
Query Match	87.2%	Score 612.4;	DB 22;	Length 1768;	
Best Local Similarity	92.0%	Pred. No. 7e-175;			
Matches	646;	Conservative 0;	Mismatches 56;	Indels 0;	Gaps 0;
Qy	1 ATGACGTATCCAAGGAGGGCGTTACCGAAGAGAACACCCGCCACCGTACCTTGCC	60			
Db	1735 ATGACATATCCAAGGAGGGCGTTACCGCAGAAGAACACCCGCCACCGTACCTTGCC	1676			
Qy	61 CAGATCCTCGCCCCCTGGCTCGTCACCCGCCACCGTACCGCTGGAGAAGG	120			
Db	1675 CAGATCCTCGCCCCCTGGCTCGTCACCCGCCACCGTACCGCTGGAGAAGG	1616			
Qy	121 AAAATGGCATCTTCAACACCCGCCCTGGATATACTGTCAGCGAAC	180			
Db	1615 AAAATGGCATCTTCAACACCCGCCCTGGATATACTGTCAGCGAAC	1556			
Qy	181 ACAGTCAGAACGCCCTCTGGGGACATGATGAGATTCAATATACTTCTT	240			
Db	1555 ACAGTCACAAACGCCCTCTGGGGACATGATGAGATTAAATTGACGACTTGT	1496			
Qy	241 CCCCAGGGGGTCAAACCCCGCTGTGCCCTTGAATACTACAGAATAAGAAAG	300			
Db	1495 CCCCGGGAGGGGACCAACAAATCTCTATACCCCTTGAATACTACAGAATAAGAAAG	1436			
Qy	301 GTTAAGGTTGAATTCTGGCCCTGCTGCCGATCACCCAGGGTAGGGCTCC	360			
Db	1435 GTTAAGGTTGAATTCTGGCCCTGCTGCCGATCACCCAGGGTAGGGCTCC	1376			
Query Match	84.8%	Score 595.6;	DB 20;	Length 1786;	

Db	1702	GTAAAGGTTGAATTCTGGCCCTGCTCCCCATCACCCAGGGTATAGGGAGTGGGCTCC	1761	CC sequalee associated with PCV-2. The present sequence is a DNA fragment of a plasmid, which expresses gene products of PCV-2.
Qy	361	AGTGCCTTATTAGATGATAACTTGTAAAGGCCAACGCCCTCACCTATGACCCC	420	CC
Db	1762	ACTGCTTATTCTAGATGATAACTTGTAAAGGCCAACGCCCTAACCTATGACCCA	1821	XX
Qy	421	TATGTAACACTCTCCGCCATACCATAACCCAGCCCTTCTCCTACCACTCCC GGTA	480	SQ Sequence 2769 BP; 828 A; 455 C; 538 G; 948 T; 0 other;
Db	1822	TATGTAACACTCTCCGCCATACCATAACCCAGCCCTTCTCCTACCACTCCC GGTA	1881	Query Match 88.6%; Score 622; DB 22; Length 2769;
Qy	481	TTTACCCCCAACCTGCTTAGATTCACTATTGATTACTTCCAACCAACAAAGA	540	Best Local Similarity 92.9%; Pred. No. 1.1e-17;
Db	1882	TTCACACCCAAACCTGTTCTGACTCCACATTGATTACTTCCAACCAATACAAAGG	1941	Matches 652; Conservative 0; Mismatches 50; Indels 0; Gaps 0;
Qy	541	AACCAAGCTGTGGCTGAGACTACAAACACTGCTGGAAATGTAGACCACGTAGGCCTGGCACT	600	1 ATGACGTATCCAAGGAGGGTTACCGAAGAAGAACCGCCCGCAGGCCATCTGGC 60
Db	1942	AATCAGCTTGGCTGAGACTACAAACCTCTGGAAATGTGGACCACTAGTAGGCTCGGGGCT	2001	1898 ATGACGTATCCAAGGAGGGCTAACCGAAGAACCGCCCGCAGGCCATCTGGC 1839
Qy	601	GGCTTCGAAAACAGTATAACGACCCAGGAATACAAATATCCGTTAACCATGTTACAA	660	61 CAGATCCTCCGGCCGGCCCTGGCTCGTCACCCGGCCACCGGTACCGCTGGAGAAGG 120
Db	2002	GGGTTCGAAAACAGTAAATACGACCCAGGAATACAAATATCCGTTAACCATGTTACAA	2061	1838 CAGATCCTCCGGCCGGCCCTGGCTCGTCACCCGGCCACCGCTACCGGTGGAGAAGG 1779
Qy	661	TTCAGAGATTAACTCTAAAGGACCCCCACTTAACCTTAA 702		121 AAAATGGCATCTTCAACACCCGGCTCTCCGGACCTTCGGATATACTGTCAAGCGAAC 180
Db	2062	TTCAGAGATTAACTCTAAAGGACCCCCACTTAACCTTAA 2103		1778 AAAATGGCATCTTCAACACCCGGCTCTCCGGACCTTCGGATATACTGTCAAGCGTAC 1719
Qy				181 ACAGTCAGAACGCCCTCTCCGGGGGACATGATGAGATTAACTTAACTGACTTTCTT 240
Db				1718 ACAGTCACAAAGGCCCTCCTGGGGGGGACATGATGAGATTAAATGACGACTTTGT 1659
Qy				241 CCCCCAGGAGGGGGTCAAACCCCCCGGCTCTGTGCCCTTGAATACTACAGAAATAAGAAAG 300
Db				1658 CCCCCGGAGGGGGACCAACAAATCTCATATACCCCTTGAATACTACAGAAATAAGAAAG 1599
Qy				301 GTTAAGGTTGAATTCTGGCCCTGCTCCCCGATCACCCAGGGTACAGGGGACTGGGCTCC 360
Db				1598 GTTAAGGTTGAATTCTGGCCCTGCTCCCCGATCACCCAGGGTACAGGGGACTGGGCTCC 1539
Qy				361 AGTGCCTGTTATTTAGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGACCCC 420
Db				1538 ACTGCTGTTATTTCTAGATGATAACTTTGTAACAAAGGCCACAGCCCTAACCTATGACCC 1479
Qy				421 TTTACCCCCAAACCTGCTCCTCCGCCATACCCAGGCCACAGCCCTCACCTATGACCCC 480
Db				1478 TATGTAACACTACTCTCCCTAACCTAACATCCCTAACCTAACATCCCTAACCTAAC 1419
RESULT 3				QY
AAF75830/C				541 AACAGCTGTGGCTGAGACTACAAACTGCTGGAAATGTAGACCACGTAGGGCTCGCAC 600
ID	AAF75830	standard; DNA; 2769 BP.		1358 ATCAGCTTGGCTGAGACTACAAACCTGTTCTGACTCCACTATTGATTACTTCCAAACCTAAC 1299
XX				601 GCCTTCGAAAACAGTATAACGACCCAGGAATACAAATATCCGCTAACCTATGACCCC 660
AC	AAF75830;			1298 GCCTTCGAAAACAGTAAATACGACCCAGGAATACAAATACAAACCTAAC 1239
XX				QY
DT	18-MAY-2001	(first entry)		661 TTCAGAGATTAACTCTAAAGGACCCCCACTTAACCTTAA 702
XX				Db
DE	pJP107	DNA fragment.		1238 TTCAGAGATTAACTCTAAAGGACCCCCACTTAACCTTAA 1197
XX				RESULT 4
KW	ALVAC; porcine circovirus-2; vaccine; pig; myocarditis; abortion;			AAF28320/C
KW	intrauterine infection; multisystemic wasting syndrome; ds.			ID AAF28320 standard; DNA; 3609 BP.
XX				XX
OS	Unidentified.			AC AAF28320;
XX				XX
PN	WO200116330-A2.			DT 30-MAR-2001 (first entry)
XX				DE PJP107 donor plasmid for PCV2 ORF2 and ORF1.
PD	08-MAR-2001.			PS Example 3 #3; Fig 6 #1; 134pp; English.
XX				XX PCV2; porcine circovirus 2; virucide; immunostimulant; vaccine;
PF	28-AUG-2000; 2000WO-EP08781.			CC postweaning multisystemic wasting syndrome; PMWS; infection;
XX				CC pig pathogen; open reading frame; ORF; ss.
PR	31-AUG-1999; 99US-0151564.			XX Porcine circovirus type 2.
XX				OS
PR	31-MAY-2000; 2000US-0583350.			
PA (MERI-)	MERIAL.			
PA (UYSA-)	UNIV SASKATCHEWAN.			
PA (UYBE-)	UNIV QUEENS BELFAST.			
XX				
PI	Ellis JA, Allan GM, Meehan B, Clark E, Haines D, Hassard L, Audonnet JP, Mcneilly F;			
XX				
DR	WPI; 2001-244408/25.			
XX				
DR	P-PSDB; AAB73273.			
XX				
PT	Use of porcine circovirus-2 immunogen to formulate a vaccine			
PT	composition to treat pigs against myocarditis, abortion, intrauterine			
PT	infection and/or post-weaning, multisystemic wasting syndrome			
PT	associated with PCV-2			
XX				
PS	Example 3 #3; Fig 6 #1; 134pp;			
XX				
CC	The present invention relates to the use of porcine circovirus-2 (PCV-2)			
CC	immunogen to formulate a vaccine composition to prevent or treat pigs			
CC	against myocarditis and/or abortion and/or intrauterine infection and/or			
CC	post-weaning, multisystemic wasting syndrome and other pathological			





DR	WPI; 2001-244408/25.	AAF28317
DR	P-PSDB; AAB73272.	ID AAF28317 standard; DNA; 2520 BP.
XX		XX
PT	Use of porcine circovirus-2 immunogen to formulate a vaccine	AC AAF28317;
PT	composition to treat pigs against myocarditis, abortion, intrauterine	XX
PT	infection and/or post-weaning, multisystemic wasting syndrome	DT 30-MAR-2001 (first entry)
PT	associated with PCV-2	XX
XX		DE pJP102 donor plasmid for PCV2 ORF2.
PS	Example 2 #3; Fig 3 #1; 134pp; English.	XX
XX	The present invention relates to the use of porcine circovirus-2 (PCV-2) immunogen to formulate a vaccine composition to prevent or treat pigs against myocarditis and/or abortion and/or intrauterine infection and/or post-weaning, multisystemic wasting syndrome and other pathological sequelae associated with PCV-2. The present sequence is a DNA fragment of a plasmid, which expresses gene products of PCV-2.	KW PCV2; porcine circovirus 2; virucide; immunostimulant; vaccine; postweaning multisystemic wasting syndrome; PMWS; infection; pig pathogen; open reading frame 2; ORF2; ss.
XX		XX
SO	Sequence 2520 BP; 777 A; 482 C; 382 G; 879 T; 0 other;	OS Porcine circovirus type 2.
SO		OS Synthetic.
SO		XX
SO		PN WO2000077216-A2.
SO		XX
SO		PD 21-DEC-2000.
SO		XX
SO		PF 09-JUN-2000; 2000WO-IB00882.
SO		XX
SO		PR 10-JUN-1999; 99US-0138478.
SO		XX
SO		PR 01-JUN-2000; 2000US-0583545.
SO		XX
SO		PA (MERI-) MERIAL.
SO		XX
SO		PI Bublot M, Perez JM, Charreyre CE;
SO		XX
SO		DR WPI; 2001-080692/09.
SO		DR P-PSDB; AAB61154.
SO		XX
SO		PT Novel recombinant virus comprising DNA from porcine circovirus 2 useful as vaccine for treatment and prophylaxis of porcine circovirus infection, such as postweaning multisystemic wasting syndrome in pigs.
SO		PT
Db	1402 ATGACGTATCCAAGGAGGGCTTACCGAAGAACACCGCATCTTGGGC	PT
Qy	1 ATGACGTATCCAAGGAGGGCTTACCGAAGAACACCGCATCTTGGGC	60
Db	1402 ATGACGTATCCAAGGAGGGCTTACCGAAGAACACCGCATCTTGGGC	1461
Qy	61 CAGATCCTCCGCCGCCCTGGCTCGTCCACCCCCGCCACCGTTACCGCTGGGAAAGC	120
Db	1462 CAGATCCTCCGCCGCCCTGGCTCGTCCACCCCCGCCACCGCTACCGCTGGGAAAGC	1521
Qy	121 AAAATGGCATCTTCAACACCCGCCCTCCCGCACCTTCGGATTAACGGCAACC	180
Db	1522 AAAATGGCATCTTCAACACCCGCCCTCCCGCACCTTCGGATTAACGGTACCC	1581
Qy	181 ACAGTCAGAACGCCCTCTGGGGGACATGATGAGATTCAATTAATGACTTTCTT	240
Db	1582 ACAGTCACAAACGCCCTCTGGGGGACATGATGAGATTAAATTGACGACTTTGTT	1641
Qy	241 CCCCAGGGGGGGTCAAACCCCCGGCTCTGTGAACTACAGATAAGAAAG	300
Db	1642 CCCCGGGAGGGGACCAACAAATCTCTATACCCCTTGAAATAAGAAAG	1701
Qy	301 GTAAAGGTTGAATTCTGGCCCTGCTCCCGATACCCAGGGGTGACAGGGGATGGCTCC	360
Db	1702 GTAAAGGTTGAATTCTGGCCCTGCTCCCGATACCCAGGGGTGATAGGGATGGCTCC	1761
Qy	361 AGTGTGTTATTAGTGTAAACTTTGTAACAAAGGCCACAGGCCCTCACCATGACCCC	420
Db	1762 ACTGTGTATTCTAGTGTAAACTTTGTAACAAAGGCCACAGGCCCTAACCTATGACCCA	1821
Qy	421 TATGTAACACTACTCTCCGGCATACCATAAACCCGGCTTACCATGACCCC	480
Db	1822 TATGTAACACTACTCTCCGGCATACCATCCCGTACCATGACCCC	1881
Qy	481 TTACCCAAACCTGTCTAGATTTCACATTGATTAACCTTCCACAAACAAAGA	540
Db	1882 TTACACCCAAACCTGTCTTGACTCCACATTGATTAACCAACAAAGG	1941
Qy	541 AACAGGTGAAACAGTATAACGAGGAAATGTAGACCTAGGGCTCGGCAC	600
Db	1942 AACAGGTGAAACAGTATAACGAGGAAATGTAGACCTAGGGCTCGGCAC	2001
Qy	601 GCGTTGAAACAGTATAACGAGGAAATACGAGGAAATCTGGTAACTATGACCAA	660
Db	2002 GCGTTGAAACAGTATAACGAGGAAATGTGGACCTACAAATAACCCCTAA	2061
Qy	661 TTCAAGAAATTAAAGACCCCCACTTAACCTTAA	702
Db	2062 TTCAAGAAATTAAAGACCCCCACTTAACCCCTAA	2103
Qy	181 ACAGTCAGAACGCCCTCTGAACTACAGAAATAAGAAAG	300
Db	1582 ACAGTCACACGCCCTCTGGGGACTACAAATAACCCCTAA	1641
Qy	241 CCCAGGGGGGGTCAAACCCCCGGCTCTGTGCCCTTGAAATAAGAAAG	300
Db	1642 CCCCGGGAGGGGGACCAACAAATCTCTATAACCCCTTGAAATAAGAAAG	1701
Qy	301 GTAAAGGTTGAATTCTGGCCCTGCTCCCCGATACCCAGGGTGACAGGGAGGGCTCC	360

NUMBER OF SEQ ID NOS: 6  
 SOFTWARE: PatentIn ver. 2.0  
 SEQ ID NO 6  
 LENGTH: 1768  
 TYPE: DNA  
 FEATURE:  
 ORGANISM: Porcine circovirus  
 NAME/KEY: variation  
 LOCATION: (1)..(1768)  
 OTHER INFORMATION: N represents A or C or G or T  
 US-09-082-558-6

Query Match 48.3%; Score 339; DB 4; Length 1768;  
 Best Local Similarity 92.2%; Pred. No. 9.4e-96;  
 Matches 357; Conservative 0; Mismatches 30; Indels 0; Gaps 0;

Qy 316 TGGCCCTGCTCCCCGATCACCCAGGGTACAGGGGAGTGGGCTCCAGTGCTGTTATTAA 375  
 Db 1768 TGGCCCTGCTCCCCCATCACCCAGGGTATAAGGGAGTGGGCTCCACTGCTGTTATTCA 1709

Qy 376 GATGATAACTTTGTAACAAGGCCACAGCCCTCACCTATGACCCCTATGTAACACTCC 435  
 Db 1708 GATGATAACTTTGTAACAAGGCCACAGCCCTAACCTATGACCCATATGTAACACTCC 1649

Qy 436 TCCCAGCCATACCCATAACCCAGCCCTTCTCCCTACCCATTGACCCATATGTAACACTCC 495  
 Db 1648 TCCCAGCCATACCAATCCCCAACCCCTTCTCCCTACCCATTGACCCATATGTAACACTCC 1589

Qy 496 GTCCTAGATTCACTATTGATTACTTCCAAACCAACAAAGAAACCCAGCTGGCTG 555  
 Db 1588 GTTCTTGACTCCACTATTGATTACTTCCAAACCCATTCTACCCATTGACCCATATGTAACACTCC 1529

Qy 556 AGACTACAAACTGCTGAAATGTAGACCACGTAGGCCCTCGGCACACTGCCTGC 615  
 Db 1528 AGGCTACAAACCTCTAGAAATGTGGACCACGTAGGCCCTCGGCACACTGCCTGC 1469

Qy 616 ATATAGCAGGAATAACATAATCCGGTACTTACCCGGTACTTACCCCAAAACT 675  
 Db 1468 ATATAGCAGGAATAACATAATCCGGTACTTACCCGGTACTTACCCCAAAACT 1409

Qy 676 TTAAAGACCCCCACTTAACCCCTAA 702  
 Db 1408 TTAAAGACCCCCACTTAACCCCTAA 1382

RESULT 15  
 US-09-267-177-24  
 ; Sequence 24, Application US/09267177  
 ; Patent No. 6287856  
 ; GENERAL INFORMATION:  
 ; APPLICANT: Poet, Steven E.  
 ; ATTORNEY: Ritchie, Branson W.  
 ; ATTORNEY: Niagro, Frank D.  
 ; ATTORNEY: Lukert, Phil D.  
 ; TITLE OF INVENTION: Vaccines against Circovirus Infections  
 ; FILE REFERENCE: 21099.0057  
 ; CURRENT APPLICATION NUMBER: US/09/267,177  
 ; CURRENT FILING DATE: 1999-03-12  
 ; EARLIER APPLICATION NUMBER: 60/077,890  
 ; EARLIER FILING DATE: 1998-03-13  
 ; NUMBER OF SEQ ID NOS: 41  
 ; SOFTWARE: FastSEQ for Windows Version 3.0  
 ; SEQ ID NO 24  
 ; LENGTH: 699  
 ; TYPE: DNA  
 ; ORGANISM: porcine circovirus  
 US-09-267-177-24

RESULT 14  
 US-09-161-092-6/c  
 ; Sequence 6, Application US/09161092  
 ; Patent No. 6391314  
 ; GENERAL INFORMATION:  
 ; APPLICANT: ALLAN, Gordon  
 ; APPLICANT: MEEHAN, Brian  
 ; APPLICANT: CLARK, Edward  
 ; APPLICANT: HAINES, Deborah  
 ; APPLICANT: HASSARD, Lori  
 ; APPLICANT: HARDING, John  
 ; APPLICANT: CHARREYRE, Catherine E.  
 ; APPLICANT: CHAPPUIS, Gilles E.  
 ; TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC  
 ; TITLE OF INVENTION: REAGENTS  
 ; FILE REFERENCE: ALLAN  
 ; CURRENT APPLICATION NUMBER: US/09/161,092  
 ; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/082,558  
 ; PRIOR FILING DATE: EARLIER FILING DATE: 1998-05-21  
 ; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: FR 9800873  
 ; PRIOR FILING DATE: EARLIER FILING DATE: 1998-01-22  
 ; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: FR 9803707  
 ; PRIOR FILING DATE: EARLIER FILING DATE: 1998-03-20  
 ; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: FR 97/12382  
 ; PRIOR FILING DATE: EARLIER FILING DATE: 1997-10-03  
 ; NUMBER OF SEQ ID NOS: 6  
 ; SOFTWARE: PatentIn ver. 2.0

Query Match 43.8%; Score 307.8; DB 4; Length 699;  
 Best Local Similarity 67.0%; Pred. No. 3e-86;  
 Matches 470; Conservative 0; Mismatches 222; Indels 9; Gaps 2;

Qy 1 ATGACGTATCCAAGGAGGCGTACCGAAGAAGACACGCCATCTGGC 60  
 Db 1 ATGACGTGGCCAAGGAGGCGTACCGAAGAAGACACGCCATCTGGC 60

Qy 61 CAGATCCTCCGCCCGCCCTGGCTCGGCTACCGCTACCGCTGG 114  
 Db 61 AACATCCTCCGGAGAACCATATTGGCACACCCCGCTTCAAAACCGTTACAGATGG 120  
 Qy 115 AGAAGGAAAATGGCATCTCAACACCCGCCTCTCCGCACCTTCGGATATACTGTCAAAG 174  
 Db 121 CGCCGAAAGACGGTATCTCAATTCCGGCTTCTACAGAATTGGTACTACCCATAAAA 180  
 Qy 175 CGAACACAGTCAGAACGCCCTCCCTGGGGGGGACATGATGAGATTCAATTAATGAC 234  
 Db 181 GGA---GGATACTCGCAGCCATCTGGATGTTAACCTCAACATCGGCCAG 237  
 Qy 235 TTCTTCCCCAGGGGGGTCAAACCCCCGGCTCTGTGCCCTTGAATACAGAATA 294  
 Db 238 TTCCCTCCCCCTCAAGGGGCAACCAACCCCTACCCCTACCTTCAAAATACCTACCGTATT 297  
 Qy 295 AGAAAGTTAAGGTGAATTCTGGCCCTGGCTCCCCGATCACCCAGGGTGACAGGGAGTG 354  
 Db 298 AGAAAGGCTAAATATGAATTTCACCCCAAGAGACCCATCACCTTAATCAAAGAGGTGTT 357  
 Qy 355 GGCTCCAGTGCTGTTATTTAGTATACTTTGTAACAAAGGCCACGCCCTCACCTAT 414  
 Db 358 GGTCACACTGTTATCTGGATGCCAACTTGTAAACCCCTCCACCCATACCTACCCCT 417  
 Qy 415 GACCCCTATGTAACACTACTCTCCGCCATACCAAACTTGTAAACAAAGGCCACGCCCTCACCTCC 474  
 Db 418 GACCCCTATATAACTACTCTCCGCCACCCATAAAGGCAGCCCTTACCTACCCCT 477  
 Qy 475 CGGTACTTTACCCCCAACCTGTCCTAGATTCACTATTGATTACTTCCAAACCAACAC 534  
 Db 478 AGGTACTTCACCCCCAACCTGAGCTGGACCAAACAAATTGATTGGTTCCACCCAAATAT 537  
 Qy 535 AAAAGAAACCAAGCTGGCTGAGACTACAAACTGCTGGAAATGTAACCCAGCTGGCCCTC 594  
 Db 538 AAAAGAAACCAAGCTGGCTCATTAAATACCCACCAATGTCGAGCCACAGGGCTC 597  
 Qy 595 GGCACCTGCGTTGAAAAACAGTATATACTGGACCCAGGAATACAAATTCGGTGAACCATGTTAT 654  
 Db 598 GGCTATGGCTCCAAATGCAAGCCACAGCCAAATATGTGGTAAGGGCTGACTATTAT 657  
 Qy 655 GTACAATTCAAGAAATTAAATTAAAGGACCCCCAACCTTAA 695  
 Db 658 GTACAATTCAAGAAATTATCTCAAAAGACCCCTCTAAATAAA 698

Search completed: May 18, 2003, 12:47:57  
 Job time : 65 secs

Db 1408 TCTTAAAGACCCCACTTAAACCTAA 1381

RESULT 9

US-09-082-558-3/C

; Sequence 3, Application US/09082558A

; Patent No. 6368601

; GENERAL INFORMATION:

; APPLICANT: ALLAN, Gordon

; APPLICANT: MEEHAN, Brian

; APPLICANT: CLARK, Edward

; APPLICANT: HAINES, Deborah

; APPLICANT: HASSARD, Lori

; APPLICANT: HARDING, John

; APPLICANT: CHARREYRE, Catherine E.

; APPLICANT: CHAPPUIS, Gilles E.

; TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC

; TITLE OF INVENTION: REAGENTS

; CURRENT APPLICATION NUMBER: US/09/082,558A

; CURRENT FILING DATE: 1998-05-21

; EARLIER APPLICATION NUMBER: FR 9800873

; EARLIER FILING DATE: 1998-01-22

; EARLIER APPLICATION NUMBER: FR 9803707

; EARLIER FILING DATE: 1998-03-20

; EARLIER APPLICATION NUMBER: FR 97/12382

; EARLIER FILING DATE: 1997-10-03

; NUMBER OF SEQ ID NOS: 6

; SOFTWARE: PatentIn Ver. 2.0

; SEQ ID NO 3

; LENGTH: 1768

; TYPE: DNA

; ORGANISM: Porcine circovirus

; US-09-082-558-3

Query Match 48.4%; Score 340; DB 4; Length 1768;

Best Local Similarity 92.3%; Pred. No. 4.6e-96; Mismatches 0; Indels 0; Gaps 0;

Matches 358; Conservative 0; Mismatches 30; Indels 0; Gaps 0;

Qy

315 CTGGCCCTGCTCCCGATACCCAGGGTACAGGGGAGTGGGACTCCAGTGTGTTATTCT 374

Db 1768 CTGGCCCTGCTCCCGATACCCAGGGTACAGGGGAGTGGGACTCCAGTGTGTTATTCT 1709

Qy

375 AGATGATAACTTTGTAACAAAGGCCACAGGCCCTCACCTATGACCCCTATGTAAACTACTC 434

Db 1708 AGATGATAACTTTGTAACAAAGGCCACAGGCCCTAACCTATGACCCATATGTAAACTACTC 1.649

Qy

435 CTCCCGCCATACCATACCCAGCCCTTCTCTTACCACTTCCGGTACTTTACCCCGAAAC 494

Db 1648 CTCCCGCCATACCATCCCGCTTCTCTTACCACTTCCGGTACTTTACCCCGAAAC 1589

Qy

495 TGTCTCTAGATTTCACATTGATTTGATACCTTCCAAACCAACAAAGAACCCAGCTGGCT 554

Db 1588 TGTTCTCTGACTCCACTATTGATTTGATACCTTCCAAACCAAAATAACAAAGGAATCAGCTTTGGCT 1529

Qy

555 GAGACTACAAACTGCTGGAAATGTAGACCCAGCTGGCCTCGGCACACTGGCTTCCGAAACAG 614

Db 1528 GAGACTACAAACCTCTGGAAATGTGGACCACTGGCTCGGGCTCGCTTGAAACAG 1469

Qy

615 TATATACGACCCAGGATAACAATATCCGTGTAACCATGTCATAATTGAAATTAA 674

Db 1468 TAAATACGACCCAGGACTACAATATCCGTGTAACCATGTCATAATTGAAATTAA 1409

Qy

675 TTTAAACACCCCCCACTTAACCCCTAA 702

Db 1408 TCTTAAAGACCCCCCACTTAACCCCTAA 1381

Qy

702 TTTAAACACCCCCCACTTAACCCCTAA 702

Db 1468 TATATACGACCCAGGACTACAATATCCGTGTAACCATGTCATAATTGAAATTAA 1409

Qy

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Db 1528 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

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Db 1528 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Qy

925 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1588 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

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982 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1528 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

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1039 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1588 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Qy

1096 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1528 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

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1153 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1588 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

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Db 1588 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

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1324 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

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1381 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

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1438 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1528 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

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1495 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1588 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

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1552 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1528 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

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1609 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1588 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

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1666 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1528 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Qy

1723 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1588 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Qy

1780 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1528 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Qy

1837 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1588 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Qy

1894 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1528 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Qy

1951 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1588 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Qy

2008 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1528 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Qy

2065 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1588 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Qy

2122 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1528 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Qy

2179 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1588 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Qy

2236 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1528 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Qy

2293 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1588 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Qy

2350 TATATACGACCCAGGATAACAACAAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Db 1528 TATATACGACCCAGGATAACAACAAACAAAGAACCCAGCTGGCTTCCGAAACAG 1469

Qy

2407 TATATACGACCCAGGATAACA

; APPLICANT: CHAPPUIS, Gilles E.  
 ; TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC  
 ; FILE REFERENCE: REAGENTS  
 ; FILE REFERENCE: ALLAN  
 ; CURRENT APPLICATION NUMBER: US/09/161,092  
 ; CURRENT FILING DATE: 1998-09-25  
 ; PRIORITY APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/082,558  
 ; PRIORITY FILING DATE: EARLIER FILING DATE: 1998-05-21  
 ; PRIORITY APPLICATION NUMBER: EARLIER APPLICATION NUMBER: FR 9800873  
 ; PRIORITY FILING DATE: EARLIER FILING DATE: 1998-01-22  
 ; PRIORITY APPLICATION NUMBER: EARLIER APPLICATION NUMBER: FR 9803707  
 ; PRIORITY FILING DATE: EARLIER FILING DATE: 1998-03-20  
 ; PRIORITY APPLICATION NUMBER: EARLIER APPLICATION NUMBER: FR 97/12382  
 ; PRIORITY FILING DATE: EARLIER FILING DATE: 1997-10-03  
 ; NUMBER OF SEQ ID NOS: 6  
 ; SOFTWARE: PatentIn Ver. 2.0  
 ; SEQ ID NO: 3  
 ; LENGTH: 1768  
 ; TYPE: DNA  
 ; ORGANISM: Porcine circovirus  
 ; US-09-161-092-4

Query Match 48.4%; Score 340; DB 4; Length 1768;  
 Best Local Similarity 92.3%; Pred. No. 4.6e-96;  
 Matches 358; Conservative 0; Mismatches 30; Indels 0; Gaps 0;

QY 315 CTGGCCCTGCTCCCCGATACCCAGGGTGCAGGGGAGTGGGCTCCAGTGCTGTATT 374  
 Db 1768 CTGGCCCTGCTCCCCCATCACCCAGGGTGTAGGGGAGTGGGCTCCACTGCTGTATTCT 1709

QY 375 AGATGATAACTTTGTAAAGGCCACAGGCCCTACCATATGACCCCTATGAACTACTC 434  
 Db 1708 AGATGATAACTTTGTAAAGGCCACAGGCCCTAACCTATGACCCATATGAACTACTC 1649

QY 435 CTCCCGCCATACCAAAACCCAGGCCCTCTCCTACCAACTCCGGTACTTTACCCCCAAC 494  
 Db 1648 CTCCCGCCATACCAATCCCCAACCCCTCTCCTACCAACTCCGGTACTTCACACCCAAAC 1589

QY 495 TGTCCTAGATTCACTTGTATGATTAACCCAAACAAACAAAGAAACAGCTGTGGCT 554  
 Db 1588 TGTTCTGTGACTCCACTATTGTAACTTCCCTACCCCTAACCTATGACCCCTATGAACTACTC 1529

QY 555 GAGACTACAAACTGCTGGAAATGTAGACCACTGGGCACCTGGGACTCGGCTCGAAACAG 614  
 Db 1528 GAGACTACAAACCTCTGGAAATGTGGACCCAGTGGCTCGGCTCGGGCTCGAAACAG 1469

QY 615 TATATAGGACCCAGGAAATACAATATCCGTGTAAACCATTGACATTGAGAAATTAA 674  
 Db 1468 TAAATAGGACCCAGGACTACAATATCCGTGTAAACCATTGACATTGAGAAATTAA 1409

QY 675 TTTAAAGACCCCCCACTTAACCCCTTA 702  
 Db 1408 TCTTAAAGACCCCCCACTTAACCCCTAA 1381

RESULT 13  
 US-09-082-558-6/C  
 ; Sequence 6, Application US/09082558A  
 ; Patent No. 6368601

; GENERAL INFORMATION:  
 ; APPLICANT: ALLAN, Gordon  
 ; APPLICANT: MEEHAN, Brian  
 ; APPLICANT: CLARK, Edward  
 ; APPLICANT: HAINES, Deborah  
 ; APPLICANT: HASSARD, Lori  
 ; APPLICANT: HARDING, John  
 ; APPLICANT: CHARREYRE, Catherine E.  
 ; APPLICANT: CHAPPUIS, Gilles E.  
 ; TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC  
 ; FILE REFERENCE: REAGENTS  
 ; CURRENT APPLICATION NUMBER: US/09/082,558A  
 ; CURRENT FILING DATE: 1998-05-21  
 ; EARLIER APPLICATION NUMBER: FR 9800873  
 ; EARLIER FILING DATE: 1998-01-22  
 ; EARLIER APPLICATION NUMBER: FR 9803707  
 ; EARLIER FILING DATE: 1998-03-20  
 ; EARLIER APPLICATION NUMBER: FR 97/12382  
 ; EARLIER FILING DATE: 1997-10-03

RESULT 12  
 US-09-161-092-4/C  
 ; Sequence 4, Application US/09161092  
 ; Patent No. 6391314

; GENERAL INFORMATION:  
 ; APPLICANT: ALLAN, Gordon  
 ; APPLICANT: MEEHAN, Brian  
 ; APPLICANT: CLARK, Edward  
 ; APPLICANT: HAINES, Deborah  
 ; APPLICANT: HASSARD, Lori  
 ; APPLICANT: HARDING, John  
 ; APPLICANT: CHARREYRE, Catherine E.  
 ; APPLICANT: CHAPPUIS, Gilles E.  
 ; TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC  
 ; FILE REFERENCE: REAGENTS  
 ; CURRENT APPLICATION NUMBER: US/09/082,558A  
 ; CURRENT FILING DATE: 1998-05-21  
 ; EARLIER APPLICATION NUMBER: FR 9800873  
 ; EARLIER FILING DATE: 1998-01-22  
 ; EARLIER APPLICATION NUMBER: FR 9803707  
 ; EARLIER FILING DATE: 1998-03-20  
 ; EARLIER APPLICATION NUMBER: FR 97/12382  
 ; EARLIER FILING DATE: 1997-10-03

APPLICANT: ELLIS, John A.  
 APPLICANT: KRAKOWKA, George S.  
 APPLICANT: AUDONNET, Jean-Christophe F.  
 TITLE OF INVENTION: PORCINE CIRCOVIRUS AND PARVOVIRUS VACCINE  
 FILE REFERENCE: 4 54 313-23 38  
 CURRENT APPLICATION NUMBER: US/09/347, 594  
 CURRENT FILING DATE: 1999-07-01  
 EARLIER APPLICATION NUMBER: 98 08777  
 EARLIER FILING DATE: 1998-07-06  
 NUMBER OF SEQ ID NOS: 5  
 SOFTWARE: PatentIn Ver. 2.1  
 SEQ ID NO 2  
 LENGTH: 1767  
 TYPE: DNA  
 ORGANISM: Porcine circovirus  
 US-09-347-594-2

Query Match 53.98; Score 378.4; DB 4; Length 1767;  
 Best Local Similarity 98.5%; Pred. No. 4.9e-108;  
 Matches 382; Conservative 0; Mismatches 6; Indels 0; Gaps 0;

Qy 315 CTGGCCCTGCTCCCGATCACCCAGGGTACAGGGGAGTGGCTCCAGTGCTGTATT 374  
 Db 1767 CTGGCCCTGCTCCCGATCACCCAGGGTACAGGGGAGTGGCTCCAGTGCTGTATTCT 1708

Qy 375 AGATGATAACTTTGTAACAAAGGCCACAGCCCTACCATATGACCCCTATGTAAACTACTC 434  
 Db 1707 AGATGATAACTTTGTAACAAAGGCCACAGCCCTACCATATGACCCCTATGTAAACTACTC 1648

Qy 375 AGATGATAACTTTGTAACAAAGGCCACAGCCCTACCATATGACCCCTATGTAAACTACTC 374  
 Db 1707 AGATGATAACTTTGTAACAAAGGCCACAGCCCTACCATATGACCCCTATGTAAACTACTC 1708

Qy 435 CTCCCGCCATACCCATACCCAGGGCTTCTCTTACCACTCCGGTACTTTACCCCTAAACCC 494  
 Db 1647 CTCCCGCCATACCCATACCCAGGGCTTCTCTTACCACTCCGGTACTTTACCCCTAAACCC 1588

Qy 495 TGTCTCTAGATTTCACATTGATTACTTCCAAACCAAAACAAAGAAACCAACTGTGGCT 554  
 Db 1587 TGTCTCTAGATTTCACATTGATTACTTCCAAACCAAAACAAAGAAATCAGCTGTGGCT 1528

Qy 555 GAGACTACAAACTGCTGGAAATGTAGACCCAGCTGGCACTTGGCTTCGAAACAG 614  
 Db 1527 GAGACTACAAACTACTGGAAATGTAGACCCAGCTGGCACTTGGCTTCGAAACAG 1468

Qy 615 TATATACGACCGGAAATACAATATCCGTGTAACCATTGTAACATTGAGAAATTAA 674  
 Db 1467 TATATACGACCGGAAATACAATATCCGTGTAACCATTGAGAAATTAA 1408

Qy 675 TTTAAAGACCCCCCACTTAACCCCTAA 702  
 Db 1407 TCTTAAGACCCCCCACTTAACCCCTAA 1380

RESULT 6  
 US-09-161-092-2/c  
 Sequence 2, Application US/09161092

Patent No. 6391314  
 GENERAL INFORMATION:  
 APPLICANT: ALLAN, Gordon  
 APPLICANT: MEEHAN, Brian  
 APPLICANT: CLARK, Edward  
 APPLICANT: HAINES, Deborah  
 APPLICANT: HASSARD, Lori  
 APPLICANT: CHARREYRE, Catherine E.  
 APPLICANT: CHAPUIS, Gilles E.  
 TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC  
 FILE REFERENCE: ALLAN  
 CURRENT APPLICATION NUMBER: US/09/082, 558A  
 CURRENT FILING DATE: 1998-05-21  
 EARLIER APPLICATION NUMBER: FR 9800873  
 PRIOR FILING DATE: EARLIER FILING DATE: 1998-01-22  
 PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: FR 9803707  
 PRIOR FILING DATE: EARLIER FILING DATE: 1998-03-20  
 PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: FR 97/12382  
 NUMBER OF SEQ ID NOS: 6  
 SOFTWARE: PatentIn Ver. 2.0  
 SEQ ID NO 2  
 LENGTH: 1767  
 TYPE: DNA

; ORGANISM: Porcine circovirus  
US-09-161-092-2

Query Match 53.9%; Score 378.4; DB 4; Length 1767;  
Best Local Similarity 98.5%; Pred. No. 4.9e-108;  
Matches 382; Conservative 0; Mismatches 6; Indels 0; Gaps 0;

Db 1648 CTCCCGCCATACAATCCCCAACCCCTCTCTTACCAACTCCCGTTACTTCACACCCAAAC 1589

Qy 495 TGTCCCTAGATTCACTTACTTGTATTCAACCAACCAACAAACAAACAAAGAAACCAACGGCTGTGGCT 554

Db 1588 TGTTCCTGACTCCACTTGTATTCAACCAAAATAACAAAGGAATCAGCTTGGCT 1529

Qy 315 CTGGCCCTGCTCCCCGATCACCCAGGGTACAGGGGAGTGGGCTCCAGTGGCTGTATT 374

Db 1767 CTGGCCCTGCTCCCCGATCACCCAGGGTACAGGGGAGTGGGCTCCAGTGGCTGTATTCT 1708

Qy 375 AGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGACCCCTATGTAAACTACTC 434

Db 1707 AGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGACCCCTATGTAAACTACTC 1648

Qy 435 CTCCGGCATACCATAACCCAGCCCTTCTTACCACTTACCCCCAAC 494

Db 1647 CTCCGGCATACCATAACCCAGCCCTTCTTACCACTTACCCCCAAC 1588

Qy 495 TGTCCCTAGATTCACTTACTTGTATTCAACCAACAAAGAAACCAACGGCTGTGGCT 554

Db 1587 TGTCCCTAGATTCCACTTACTTGTATTCAACCAACAAAGAAATCAGCTGTGGCT 1528

Qy 555 GAGACTACAAACTGCTGGAAATGTAGACCACCGTAGGGCACTCGGCTTCGCGTTTCGAAAAACAG 614

Db 1527 GAGACTACAAACTACTGGAAATGTAGACCACCGTAGGGCACTCGGCTTCGCGTTTCGAAAAACAG 1468

Qy 615 TATATACGACCCAGGAATACAATATCAGAGAATTAA 674

Db 1467 TATATACGACCCAGGAATACAATATCAGAGAATTAA 1408

Qy 675 TTTAAAGACCCCCACTTAACCCCTAA 702

Db 1407 TCTTAAAGACCCCCACTTAACCCCTAA 1380

RESULT 7  
US-09-347-594-3/C  
; Sequence 3, Application US/09347594  
; Patent No. 6217883

; GENERAL INFORMATION:  
; APPLICANT: ALLAN, Gordon M.  
; APPLICANT: MEEHAN, Brian M.  
; APPLICANT: ELLIS, John A.  
; APPLICANT: KRAKOWKA, George S.  
; APPLICANT: AUDONNET, Jean-Christophe F.  
; TITLE OF INVENTION: PORCINE CIRCOVIRUS AND PARVOVIRUS VACCINE  
; FILE REFERENCE: 454313-2338  
; CURRENT APPLICATION NUMBER: US/09/347,594  
; CURRENT FILING DATE: 1999-07-01  
; EARLIER APPLICATION NUMBER: 98 08777  
; EARLIER FILING DATE: 1998-07-06  
; NUMBER OF SEQ ID NOS: 5  
; SOFTWARE: PatentIn Ver. 2.1  
; SEQ ID NO 4  
; LENGTH: 1768  
; TYPE: DNA  
; ORGANISM: Porcine circovirus  
; US-09-347-594-4

Query Match 48.4%; Score 340; DB 4; Length 1768;  
Best Local Similarity 92.3%; Pred. No. 4.6e-96;  
Matches 358; Conservative 0; Mismatches 30; Indels 0; Gaps 0;

Db 1768 CTGGCCCTGCTCCCCATCACCCAGGGTATAGGGGACTGGCTTACTGCTGTATTCT 1709

Qy 315 CTGGCCCTGCTCCCCATCACCCAGGGTACACGGGACTGGCTTACTGCTGTATTCT 374

Db 1708 AGATGATAACTTTGTAACAAAGGCCACAGCCCTAACCTATGACCCATATGTAAACTACTC 1649

Qy 435 CTCCGGCATACCATAACCCAGCCCTTCTTACCAACTCCGGTACTTTACCCCCAAC 494

Db 1648 CTCCGGCATACCATAACCCCTCTCTTACCAACTCCGGTTACTTCACACCCAAAC 1589

Qy 495 TGTCCCTAGATTCACTTACTTGTATTCAACCAACAAACAAAGAAACCAACGGCTGTGGCT 554

Db 1588 TGTTCCTGACTCCACTTGTATTCAACCAAAATAACAAAGGAATCAGCTTGGCT 1529

Qy 555 GAGACTACAAACTGCTGGAAATGTAGACCACCGTAGGGCTCGGCTTCGCGTTTCGAAAAACAG 614

Db 1528 GAGACTACAAACCTCTGGAAATGTGGGACTAGGGGACTGGCTCGGCTTCGCGTTTCGAAAAACAG 1469

Qy 615 TATATACGACCCAGGAATACAATATCAGAGAATTAA 674

Db 1468 TAAATACGACCCAGGAATACAATATCAGAGAATTAA 1409

Qy 675 TTTAAAGACCCCCACTTAACCCCTAA 702



Db 1467 TATATACGACCGAAATAACAATATCCGTGTAACCATGATGTACAATTCAAGAGAATTAA 1408  
 QY 675 TTTAAAGACCCCACTTAAACCTTAA 702  
 Db 1407 TCTAAAGACCCCACTTAAACCTTAA 1380

RESULT 2  
 US-09-082-558-1/c  
 ; Sequence 1, Application US/09082558A  
 ; Patent No. 639134  
 ; GENERAL INFORMATION:  
 ; APPLICANT: ALLAN, Gordon  
 ; APPLICANT: MEEHAN, Brian  
 ; APPLICANT: CLARK, Edward  
 ; APPLICANT: HAINES, Deborah  
 ; APPLICANT: HASSARD, Lori  
 ; APPLICANT: HARDING, John  
 ; APPLICANT: CHARREYRE, Catherine E.  
 ; APPLICANT: CHAPPUIS, Gilles E.  
 ; TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC  
 ; FILE REFERENCE: ALLAN  
 ; CURRENT APPLICATION NUMBER: US/09/161,092  
 ; CURRENT FILING DATE: 1998-09-25  
 ; PRIORITY APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/082,558  
 ; PRIORITY FILING DATE: EARLIER FILING DATE: 1998-05-21  
 ; PRIORITY APPLICATION NUMBER: EARLIER APPLICATION NUMBER: FR 9800873  
 ; PRIORITY FILING DATE: EARLIER FILING DATE: 1998-01-22  
 ; PRIORITY APPLICATION NUMBER: EARLIER APPLICATION NUMBER: FR 9803707  
 ; PRIORITY FILING DATE: EARLIER FILING DATE: 1998-03-20  
 ; PRIORITY APPLICATION NUMBER: EARLIER APPLICATION NUMBER: FR 97/12382  
 ; PRIORITY FILING DATE: EARLIER FILING DATE: 1997-10-03  
 ; NUMBER OF SEQ ID NOS: 6  
 ; SOFTWARE: PatentIn Ver. 2.0  
 ; SEQ ID NO 1  
 ; LENGTH: 1767  
 ; TYPE: DNA  
 ; ORGANISM: Porcine circovirus

US-09-082-558-1

Query Match 54.4%; Score 381.6; DB 4; Length 1767;  
 Best Local Similarity 99.0%; Pred. No. 5e-109;  
 Matches 384; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 315 CTGGCCCTGCTCCCGATCACCCAGGGTACAGGGACTGGCTCCAGTGCTGTATT 374  
 Db 1767 CTGGCCCTGCTCCCGATCACCCAGGGTACAGGGACTGGCTCCAGTGCTGTATTCT 1708

QY 375 AGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGACCCCTATGTAAACTACTC 434  
 Db 1707 AGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGACCCCTATGTAAACTCTC 1648

QY 435 CTCCCGCCATACCATAACCCAGCCATACCCAGGGTACAGGGACTGGCTCCAGTGCTGTATTCT 1708  
 Db 1707 AGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGACCCCTATGTAAACTACTC 1648

QY 435 CTCCCGCCATACCATAACCCAGCCATACCCAGGGTACAGGGACTGGCTCCAGTGCTGTATTCT 1708  
 Db 1647 CTCCCGCCATACCATAACCCAGCCATACCCAGGGTACAGGGACTGGCTCCAGTGCTGTATTCT 1648

QY 495 TGTCCTAGATTTCACATTGATTACTTCACCTATGACCCCTATGTAAACTACTC 434  
 Db 1587 TGTCCTAGATTTCACATTGATTACTTCACCTATGACCCCTATGTAAACTACTC 1588

QY 495 TGTCCTAGATTTCACATTGATTTGAAATTGAGACCACTGGGACTGGCTCGAAACAG 494  
 Db 1587 TGTCCTAGATTTCACATTGATTTGAAATTGAGACCACTGGGACTGGCTCGAAACAG 1528

QY 555 GAGACTACAAACTGCTGGAATTGAGACCACTGGGACTGGCTCGAAACAG 614  
 Db 1527 GAGACTACAAACTGCTGGAATTGAGACCACTGGGACTGGCTCGAAACAG 1468

QY 615 TATATACGACCGAGGATAACAATATCCGTGTAACCATGATGTACAATTCAAGAGAATTAA 674  
 Db 1467 TATATACGACCGAGGATAACAATATCCGTGTAACCATGATGTACAATTCAAGAGAATTAA 1408

QY 675 TTTAAAGACCCCACTTAA 702  
 Db 1407 TCTAAAGACCCCACTTAA 1380

RESULT 4  
 US-09-347-594-2/c  
 ; Sequence 2, Application US/09347594  
 ; Patent No. 6217883  
 ; GENERAL INFORMATION:  
 ; APPLICANT: ALLAN, Gordon M.  
 ; APPLICANT: MEEHAN, Brian M.

RESULT 2  
 CNS00CNG  
 LOCUS  
 DEFINITION CNS00CNG 939 bp DNA linear GSS 04-JUN-1999  
 Drosophila melanogaster genome survey sequence TET3 end of BAC #  
 BACR26H16 of RPCI-98 library from Drosophila melanogaster (fruit  
 fly), genomic survey sequence.  
 AL059400  
 AL059400.1 GI:4946964  
 GSS.  
 SOURCE Drosophila melanogaster.  
 ORGANISM Drosophila melanogaster  
 Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;  
 Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;  
 Ephydriodea; Drosophilidae; Drosophila.  
 1 (bases 1 to 939)  
 REFERENCE Genoscope.  
 AUTHORS Direct Submission  
 JOURNAL Submitted (02-JUN-1999) Genoscope - Centre National de Sequencage :  
 BP 191 91006 EVRY cedex - FRANCE (E-mail : seqref@genoscope.cns.fr)  
 COMMENT - Web : [www.genoscope.cns.fr](http://www.genoscope.cns.fr)  
 Determination of this BAC-end sequence was carried out as part of a  
 collaboration with the Berkeley Drosophila Genome Project (BDGP).  
 The BDGP is constructing a physical map of the Drosophila  
 melanogaster genome using these BACs. For further information  
 please see <http://www.fruitfly.org> The BDGP Drosophila  
 melanogaster BAC library was prepared by Kazutoyo Osoegawa and  
 Aaron Mammoser in Pieter de Jong's laboratory in the Department of  
 Cancer Genetics at the Roswell Park Cancer Institute in Buffalo,  
 NY. The library is named RPCI-98 and was constructed by partial  
 ECO RI digestion of Drosophila DNA provided by the BDGP from the  
 isogenic strain y2; cn bw sp, the same strain used for the BDGP's  
 P1 and EST libraries. A more detailed description of the library  
 and how to order individual BAC clones, the entire library, or  
 filters for hybridization from the BACPAC Resource Center can be  
 found at [http://bacpac.med.buffalo.edu/drosophila\\_bac.htm](http://bacpac.med.buffalo.edu/drosophila_bac.htm).  
 FEATURES Location/Qualifiers  
 source  
 1. .939  
 /organism="Drosophila melanogaster"  
 /db\_xref="taxon:7227"  
 /clone="BACR26H16"  
 /clone\_lib="RPCI-98"  
 /note="end : TET3"  
 71 a 349 c. 104 g 180 t 235 others  
 BASE COUNT  
 COUNT

Query Match	6.48;	Score	44.6;	DB	17;	Length	939;
Best Local Similarity	18.18;	Pred. No.	0.23;				
Matches 56;	Conservative	122;	Mismatches	131;	Indels	0;	Gaps 0;
QY	391	ACAAAGGCCACAGCCCTCACCTATGACCCCTATGTAAACTACTCCTCCCCCAT'ACCCATA	450				
Db	274	MMMHAWMCCTTCCCTCCCYCMCYTCCCCCAYYMCCCCCCTCTYCCCCATCCMMC	33.3				
QY	451	ACCCAGGCCCTTCCACTTACCCGGTACTTTACCCCAAAACCTGTCCTAGATT'CACT	510				
Db	334	CCCTCTCCCCCMCCYATCCCTCMMMHHTTYMMCCCCCMCMMMCHMTCCMMMM	393				
QY	511	ATTGATTACTCCAACCAAACAAAGAACCAACCGCTGTGGCTGAGACTACAAACTGCT	570				
Db	394	MMMMMMMTMMHMMMMMMMTMMHMTMMMCMTMTMMTTMMTTMM	453				
QY	571	GGAAATGTTAGACCACCGTAGGCCACTGCCACTGCGAACAGTATAACGACCAGGAA	630				
Db	454	MMMMMMMMMTMMMMMMMTMMHMMMMMMHMMHMTMMMMHMMHMMHMM	513				
QY	631	TACAATATCCGTTAACCATGTTACAAATTCAAGAATTAAAGACCCCCCA	690				
Db	514	MMMMMMMTMMMMMMMTMMHMMMMMMHMMHMMMMHMMHMMHMM	573				
QY	691	CTTAACCCCT	699				
Db	574	CGTCTCTCT	582				
RESULT	3						
LOCUS	BI629442/c						
DEFINITION		638 bp mRNA linear EST 10-SEP-2001					
	RH58310.5prime RH Drosophila melanogaster normalized Head pFLC-1						
	Drosophila melanogaster cDNA clone RH58310 5 similar to CG17108:						
	FBan0017108 GO:[] located on: 2L 32A1-32A1; 08/23/2001, mRNA						
SEQUENCE		sequence.					
VERSION	BI629442.1	GI:15531652					
KEYWORDS	EST						
SOURCE	fruit fly						
ORGANISM	Drosophila melanogaster						
AUTHORS	Eukaryota; Arthropoda; Hexapoda; Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Ephydriidea; Drosophilidae; Drosophila.						
REFERENCE	1 (bases 1 to 638)						
	Stapleton, M., Brokstein, P., Hong, L., Tyler, D., Berman, B., Carlson, J., Champe, M., Chavez, C., Dorsett, V., Farfan, D., Frise, E., George, R., Gonzalez, M., Guarin, H., Harris, N., Li, P., Liao, G., Misra, S., Mungall, C.J., Nunoo, J., Pacleb, J., Paragas, V., Park, S., Phouanenavong, S., Wan, K., Yu, C., Lewis, S.E., Celniker, S. and Rubin, G.M.						
TITLE	BDGP/RH Drosophila EST Project						
JOURNAL	Unpublished (2001)						
COMMENT	Contact: Stapleton, M.						
FEATURES	BDGP						
Source	Lawrence Berkeley National Lab						
	One Cyclotron Rd, Berkeley, CA 94720, USA						
	Fax: 510 486 6798						
	Email: <a href="http://www.fruitfly.org/EST/est@fruitfly.berkeley.edu">http://www.fruitfly.org/EST/est@fruitfly.berkeley.edu</a>						
	hit genomic AE003629; arm: 2L [10413221,10674011]						
	estimated-cyto: 31E5-32A5: 08/23/2001						
	Plate: RH583 row: A column: 10						
	High quality sequence stop: 511.						
	Location/Qualifiers						
	1. 638						
	/organism="Drosophila melanogaster"						
	/db_xref="taxon:7227"						

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/clone="RH38310"
/clone_lib="RH Drosophila melanogaster normalized Head
pFLc-1"
/sex="male and female"
/dev_stage="Adult"
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GenCore version 5.1.6  
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OM nucleic - nucleic search, using sw model

Run on: May 18, 2003, 10:32:44 ; Search time 1499 Seconds

(without alignments)  
 7584.544 Million cell updates/sec

Title: US-09-514-245B-25

Perfect score: 702  
 Sequence: 1 attacggtatccaaaggaggcg.....accccccacctaaccctaa 702

Scoring table: IDENTITY\_NUC  
 Gapop 10.0 , Gapext 1.0

Searched: 16154066 seqs, 8097743376 residues

Total number of hits satisfying chosen parameters: 32308132

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : EST:\*

1: em\_estba:\*

2: em\_esthum:\*

3: em\_estin:\*

4: em\_estmu:\*

5: em\_estov:\*

6: em\_estpl:\*

7: em\_estro:\*

8: em\_htc:\*

9: gb\_est1:\*

10: gb\_est2:\*

11: gb\_htc:\*

12: gb\_est3:\*

13: gb\_est4:\*

14: gb\_est5:\*

15: em\_estfun:\*

16: em\_estom:\*

17: gb\_gss:\*

18: em\_gss\_hum:\*

19: em\_gss\_inv:\*

20: em\_gss\_pln:\*

21: em\_gss\_vrt:\*

22: em\_gss\_fun:\*

23: em\_gss\_mam:\*

24: em\_gss\_mus:\*

25: em\_gss\_other:\*

26: em\_gss\_pro:\*

27: em\_gss\_rod:\*

## ALIGNMENTS

RESULT 1  
 AI514058/c  
 LOCUS GH27112.5prime GH Drosophila melanogaster head POT2 Drosophila melanogaster cDNA clone GH27112 5prime, mRNA sequence.

DEFINITION melanogaster

ACCESSION AI514058  
 VERSION AI514058.1  
 KEYWORDS EST

SOURCE fruit fly

ORGANISM Drosophila melanogaster

Eukaryota; Metazoa; Arthropoda; Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Ephydriidae; Drosophilidae; Drosophila.

REFERENCE 1 (bases 1 to 561)  
 Harvey, D., Brokstein, P., Hong, L., Evans-Holm, M., Lewis, S., and Rubin, G.M.

AUTHORS BDGP/HM1 Drosophila EST Project

TITLE Unpublished (2001)

JOURNAL Contact: Stapleton, M.

COMMENT BDGP

Lawrence Berkeley National Lab  
 One Cyclotron Rd, Berkeley, CA 94720, USA  
 Fax: 510 486 6798  
 Email: <http://www.fruitfly.org/EST>, est@fruitfly.berkeley.edu  
 Plate: 271 row: A column: 12  
 High quality sequence stop: 488.

FEATURES Location/Qualifiers  
 1. .561  
 /organism="Drosophila melanogaster"

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
C 1	45.4	6.5	561	9 AI514058	AI514058 GH27112.5
C 2	44.6	6.4	939	17 CNS00CNG	AL059400 Drosophil
C 3	44.2	6.3	638	13 BI629442	BI629442 RH58310.5
C 4	44	6.3	445	9 AI292664	AI292664 GH15617.5
C 5	44	6.3	484	9 AI108264	AI108264 GH07058.5
C 6	44	6.3	493	9 AI405757	AI405757 GH25879.5

ACCESSION AI405757.1 GI:4248844  
 VERSION EST  
 SOURCE fruit fly.  
 ORGANISM Drosophila melanogaster  
 Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;  
 Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;  
 Ephydriodea; Drosophilidae; Drosophila.  
 1. (bases 1 to 493)  
 Harvey, D., Brokstein, P., Hong, L., Evans-Holm, M., Su, C., Tsang, G.,  
 Lewis, S., and Rubin, G.M.  
 BDGP/HHMI Drosophila EST Project  
 Unpublished (2001)  
 COMMENT Contact: Stapleton, M.

TITLE BDGP  
 JOURNAL Lawrence Berkeley National Lab  
 COMMENT One Cyclotron Rd, Berkeley, CA 94720, USA  
 FAX: 510 486 6798  
 Email: <http://www.fruitfly.org/EST>, [est@fruitfly.berkeley.edu](mailto:est@fruitfly.berkeley.edu)  
 Plate: 143 row: C column: 8  
 High quality sequence stop: 457.  
 FEATURES  
 source

1. .497  
 /organism="Drosophila melanogaster"  
 /db\_xref="taxon:7227"  
 /clone="GH14332"  
 /clone\_lib="GH Drosophila melanogaster head por2"  
 /sex="male and female"  
 /dev\_stage="adult"  
 /lab\_host="DH5 - alpha"  
 /note="Organ: head; Vector: por2; Site\_1: EcoRI; Site\_2:  
 XhoI; Sized fractionated cDNAs were directly ligated into  
 por2. Plasmid cDNA library."  
 BASE COUNT 80 a 96 c 222 g 99 t  
 ORIGIN

Query Match 6.38; Score 44; DB 9; Length 497;  
 Best Local Similarity 58.38; Pred. No. 0.29;  
 Matches 0; Mismatches 55; Indels 0; Gaps 0;

QY 35 GACACGGCCCCGGAGCCATCTGGCCAGATCTGGCTGGCCCTGGCTCGTCCAC 94  
 Db 181 GATTCCACCGCCTCACCCAGAAATGGCCACCTCGCCGATTCGGCCACCAATCCAG 122

QY 95 CCCGCCACCGTTACCGCTGGAGAAGGAAAAATGGCATCTCAACACCCGCCTCTCCCGCA 154  
 Db 121 GCCGCCACCGATTCCGGCTACCTCCACCAATGGCCACCTCCACCGGATTCGGCCAC 62

QY 155 CCTTCGGATATA 166  
 Db 61 TCCTCCGGATA 50

Query Match 6.38; Score 44; DB 9; Length 493;  
 Best Local Similarity 58.38; Pred. No. 0.29;  
 Matches 0; Mismatches 55; Indels 0; Gaps 0;

RESULT 8  
 AI134557/c  
 LOCUS AI134557  
 DEFINITION GH12051.5 prime GH Drosophila melanogaster head por2 Drosophila melanogaster cDNA clone GH12051.5 prime, mRNA sequence.  
 ACCESSION AI134557  
 VERSION AI134557.1  
 KEYWORDS EST  
 SOURCE fruit fly.  
 ORGANISM Drosophila melanogaster  
 Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;  
 Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;  
 Ephydriodea; Drosophilidae; Drosophila.  
 REFERENCE Harvey, D., Brokstein, P., Hong, L., Evans-Holm, M., Su, C., Tsang, G.,  
 Lewis, S., and Rubin, G.M.  
 TITLE BDGP/HHMI Drosophila EST Project  
 JOURNAL Unpublished (2001)  
 COMMENT 1 (bases 1 to 532)  
 BDGP

Lawrence Berkeley National Lab  
 One Cyclotron Rd, Berkeley, CA 94720, USA  
 FAX: 510 486 6798  
 Email: <http://www.fruitfly.org/EST>, [est@fruitfly.berkeley.edu](mailto:est@fruitfly.berkeley.edu)  
 Plate: 120 row: E column: 3  
 High quality sequence stop: 403.  
 FEATURES  
 source

1. .532  
 /organism="Drosophila melanogaster"  
 /db\_xref="taxon:7227"  
 /clone="GH12051"  
 /clone\_lib="GH Drosophila melanogaster head por2"  
 /sex="male and female"  
 /dev\_stage="adult"  
 /lab\_host="DH5 - alpha"

ACCESSION AI238309/c  
 VERSION AI238309.1 GI:3833167  
 SOURCE fruit fly.  
 ORGANISM Drosophila melanogaster  
 Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;  
 Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;  
 Ephydriodea; Drosophilidae; Drosophila.  
 1. (bases 1 to 497)  
 Harvey, D., Brokstein, P., Hong, L., Evans-Holm, M., Su, C., Tsang, G.,  
 Lewis, S., and Rubin, G.M.  
 BDGP/HHMI Drosophila EST Project  
 Unpublished (2001)  
 COMMENT Contact: Stapleton, M.  
 BDGP



KEYWORDS EST  
 SOURCE fruit fly:  
 ORGANISM *Drosophila melanogaster*  
 Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;  
 Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;  
 Ephydriodea; Drosophilidae; Drosophila.  
 1 (bases 1 to 58)  
 REFERENCE Harvey, D., Brokstein, P., Hong, L., Evans-Holm, M., Su, C., Tsang, G.,  
 Lewis, S., and Rubin, G.M.  
 AUTHORS BDGP/HMMI Drosophila EST Project  
 TITLE Unpublished (2001)  
 JOURNAL  
 COMMENT Contact: Stapleton, M.  
 BDGP  
 Lawrence Berkeley National Lab  
 One Cyclotron Rd, Berkeley, CA 94720, USA  
 FAX: 510 486 6798  
 Email: <http://www.fruitfly.org/EST>, [est@fruitfly.berkeley.edu](mailto:est@fruitfly.berkeley.edu)  
 Plate: 146 row: C column: 8  
 High quality sequence stop: 489.

FEATURES Source  
 1. 558 Location/Qualifiers  
 /organism="Drosophila melanogaster"  
 /db\_xref="taxon:7227"  
 /clone="GH14632"  
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 /sex="male and female"  
 /dev\_stage="adult"  
 /lab\_host="DH5 - alpha"  
 /note="Organ: head; Vector: POT2; Site\_1: EcoRI; Site\_2: XhoI; Sized fractionated cDNAs were directly ligated into POT2. Plasmid cDNA library."  
 BASE COUNT 75 a  
 ORIGIN 109 c 241 g 133 t

Query Match 6.3%; Score 44; DB 9; Length 561;  
 Best Local Similarity 58.3%; Pred. No. 0.3;  
 Matches 77; Conservative 0; Mismatches 55; Indels 0; Gaps 0;

QY 35 GACACGGCCCCGGAGCCATCTGGCCAGATCCTCGGCCAGATCCCTGGCCCTGGCTCGTCCAC 94  
 Db 558 GATTCCACCGCCTCCACCCAGAAATGGCCACCTTCCGGCAGATTCCGCCACAAATCCAGG 499

QY 95 CCCGCCACCGTTACCGCTGGAGAAGGAAATAATGGCATCTTCAACACCCGGCTCTCCCGCA 154  
 Db 498 GCGGCCACCGATTCCGCTACCTCCACCCAGAAATGGCCACCTCCACCGGATTCGGCCAC 439

QY 155 CCTTCGGATATA 166  
 Db 438 TCCTCCGGATA 427

Query Match 6.3%; Score 44; DB 9; Length 558;  
 Best Local Similarity 58.3%; Pred. No. 0.3;  
 Matches 77; Conservative 0; Mismatches 55; Indels 0; Gaps 0;

QY 35 GACACGGCCCCGGAGCCATCTGGCCAGATCCTCGGCCAGATCCCTGGCCCTGGCTCGTCCAC 94  
 Db 558 GATTCCACCGCCTCCACCCAGAAATGGCCACATCCGGCCGATTCCGCCACAAATCCAGG 499

QY 95 CCCGCCACCGTTACCGCTGGAGAAGGAAATAATGGCATCTTCAACACCCGGCTCTCCCGCA 154  
 Db 498 GCGGCCACCGATTCCGCTACCTCCACCCAGAAATGGCCACCTCCACCGGATTCGGCCAC 439

QY 155 CCTTCGGATATA 166  
 Db 438 TCCTCCGGATA 427

RESULT 12 AI404901/c  
 LOCUS GH24758\_5prime  
 DEFINITION GH Drosophila melanogaster head POT2 Drosophila  
 mRNA sequence.  
 ACCESSION AI404901  
 VERSION AI:4247988  
 KEYWORDS fruit fly:  
 SOURCE  
 ORGANISM *Drosophila melanogaster*  
 Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;  
 Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;  
 Ephydriodea; Drosophilidae; Drosophila.  
 1 (bases 1 to 561)  
 REFERENCE Harvey, D., Brokstein, P., Hong, L., Evans-Holm, M., Su, C., Tsang, G.,  
 Lewis, S., and Rubin, G.M.  
 AUTHORS BDGP/HMMI Drosophila EST Project  
 TITLE Unpublished (2001)  
 JOURNAL  
 COMMENT Contact: Stapleton, M.  
 BDGP  
 Lawrence Berkeley National Lab  
 One Cyclotron Rd, Berkeley, CA 94720, USA  
 FAX: 510 486 6798  
 Email: <http://www.fruitfly.org/EST>, [est@fruitfly.berkeley.edu](mailto:est@fruitfly.berkeley.edu)  
 Plate: 264 row: G column: 6  
 High quality sequence stop: 494.

FEATURES Source  
 1. 561 Location/Qualifiers  
 /organism="Drosophila melanogaster"  
 /db\_xref="taxon:7227"  
 /clone="GH26478"  
 /sex="male and female"  
 /dev\_stage="adult"  
 /lab\_host="DH5 - alpha"  
 /note="Organ: head; Vector: POT2; Site\_1: EcoRI; Site\_2: XhoI; Sized fractionated cDNAs were directly ligated into

/note="Organ: head; vector: pOT2; Site\_1: EcoRI; Site\_2: XbaI; Sized fractionated cDNAs were directly ligated into pOT2. Plasmid cDNA library."  
 73 a 101 c 234 g 124 t

RESULT 9  
DOCUS AI404106/c  
DEFINITION AI404106 557 bp mRNA linear EST 19-APR-2001  
GH23705.5prime GH Drosophila melanogaster head pot2 Drosophila  
melanogaster cDNA clone GH23705 5prime, mRNA sequence.  
ACCESSION AI404106  
VERSION AI404106.1 GI:4247193  
KEYWORDS EST  
SOURCE fruit fly:  
ORGANISM Drosophila melanogaster  
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;  
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;

REFERENCE	Ephydriidae; Drosophilidae; Drosophila.
AUTHORS	1 (bases 1 to 557) Harvey, D., Brokstein, P., Hong, L., Evans-Holm, M., Su, C., Tsang, G., Lewis, S. and Rubin, G. M.
TITLE	BDGP/HHMI Drosophila EST Project
JOURNAL	Unpublished (2001)
COMMENT	Contact: Stapleton, M. BDGP

FEATURES	source	High quality sequence stop: 455. Location/Qualifiers
	1. .557	/organism="Drosophila melanogaster" /db_xref="taxon:7227" /clone="GH23705" /clone_lib="GH Drosophila melanogaster head pOT2" /sex="male and female" /dev_stage="adult" /lab_host="DH5 - alpha" /note="Organ: head; Vector: pOT2; Site_1: EcoRI; Site_2: XhoI; Sized fractionated cDNAs were directly ligated into pOT2. Plasmid cDNA library." 87 3 100 3 255 3 105 +

RESULT 10  
 AI406114/c  
 LOCUS AI406114  
 DEFINITION 557 bp mRNA linear EST 19-APR-200  
 melanogaster GH26313.5prime GH Drosophila melanogaster head pOT2 Drosophila  
 melanogaster cDNA clone GH26313 5prime, mRNA sequence.  
 .  
 ACCESSION AI406114  
 VERSION AI406114.1 GI:4249201  
 KEYWORDS EST  
 SOURCE fruit fly.  
 ORGANISM Drosophila melanogaster  
 Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;  
 Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;  
 Ephydriidea; Drosophilidae; Drosophila.

AUTHORS	Harvey, D., Brokstein, P., Hong, L., Evans-Holm, M., Su, C., Tsang, G., Lewis, S. and Rubin, G.M.
TITLE	BDGP/HMM Drosophila EST Project
JOURNAL	Unpublished (2001)
COMMENT	Contact: Stapleton, M. BDGP
FEATURES	Lawrence Berkeley National Lab One Cyclotron Rd, Berkeley, CA 94720, USA Fax: 510 486 6798 Email: <a href="http://www.fruitfly.org/EST">http://www.fruitfly.org/EST</a> , est@fruitfly.berkeley.edu Plate: 263 row: B column: 1 High quality sequence stop: 494. Location/Qualifiers

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source
  .33/
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  /db_xref="taxon:7227"
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  /dev_stage="adult"
  /lab_host="DH5 - alpha"
  /note="Organ: head; Vector: pOT2; Site_1: ECORI; Site_2:
XHOI; Sized fractionated cDNAs were directly ligated into
pOT2. Plasmid cDNA library."
  74 a 109 c 242 g 132 t
  BASE COUNT

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Query	Match	6.38;	Score	44;	DB	9;	Length	557;
Best	Local	Similarity	58.38;	Pred.	NO.	0.3;		
Matches	77;	Conservative	.0;	Mismatches	55;	Indels	0;	Gaps
Qy	35	GACACGGCCCCGGAGCCATCTGGCCAGATCCTCCGGCCGGCCGGCCCTGGCTCGTCCACCC	94					
Db	557	GATTCCACCCGCCCTCCACCAATGCCACCCAGAATGCCACCCGATTCCGCCACCAATCCAGG	498					
Qy	95	CCGCCACCGTTACCGCTGGAGAAAAATGGCATCTCAACACCCGCCCTCCGGCA	154					
Db	497	GCCGCCACCGATTCCGGCTACCTCCACCAAGAATGGCCACCTCCACCCAGAATCCGCCACCC	438					
Qy	155	CCGCCACCCGATGAA	166					

RESULT 11  
 AI238565/C  
 LOCUS  
 DEFINITION 558 bp mRNA linear EST 19-APR-200  
 GH14632.5prime GH Drosophila melanogaster head pot2 Drosophila  
 melanogaster cDNA clone GH14632 5prime, mRNA sequence.  
 ACCESSION AI238565  
 VERSION AI238565.1 GI:3833423



RESULT <sup>2</sup> US-09-935-428A-1/C  
 ; Sequence 1, Application US/09935428A  
 ; Patent No. US20020106639A1  
 ; GENERAL INFORMATION:  
 ; APPLICANT: WANG, LI  
 ; APPLICANT: BABIUK, LORNE A.  
 ; APPLICANT: POTTER, ANDREW A.  
 ; APPLICANT: WILLSON, PHILIP  
 ; TITLE OF INVENTION: POSTWEANING MULTISYSTEM WASTING SYNDROME VIRUS FROM PIGS  
 ; FILE REFERENCE: 9000-0040  
 ; CURRENT APPLICATION NUMBER: US/09/935, 428A  
 ; CURRENT FILING DATE: 2001-08-20  
 ; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/209, 961  
 ; PRIOR FILING DATE: EARLIER FILING DATE: 1998-12-10  
 ; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/069, 233  
 ; PRIOR FILING DATE: EARLIER FILING DATE: 1997-12-11  
 ; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/069, 750  
 ; PRIOR FILING DATE: EARLIER FILING DATE: 1997-12-16  
 ; SEQ ID NO 1  
 ; SOFTWARE: PatentIn Ver. 2.0  
 ; LENGTH: 1768  
 ; TYPE: DNA  
 ; ORGANISM: Porcine Circovirus Type II  
 ; US-09-935-428A-1

Query Match 88.4%; Score 620.4; DB 10; Length 1768;  
 Best Local Similarity 92.7%; Pred. No. 2.3e-195;  
 Matches 651; Conservative 0; Mismatches 51; Indels 0; Gaps 0;

1 ATGACGTATCCAAGGAGGGCTTACCGAAGAGAACCCGGCAGGCATCTTGGC 60  
 |||||||

RESULT <sup>3</sup> US-09-935-428A-24/C  
 ; Sequence 24, Application US/09935428A  
 ; Patent No. US20020106639A1  
 ; GENERAL INFORMATION:  
 ; APPLICANT: WANG, LI  
 ; APPLICANT: BABIUK, LORNE A.  
 ; APPLICANT: POTTER, ANDREW A.  
 ; APPLICANT: WILLSON, PHILIP  
 ; TITLE OF INVENTION: POSTWEANING MULTISYSTEM WASTING SYNDROME VIRUS FROM PIGS  
 ; FILE REFERENCE: 9000-0040  
 ; CURRENT APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/209, 961  
 ; CURRENT FILING DATE: EARLIER FILING DATE: 1998-12-10  
 ; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/069, 233  
 ; PRIOR FILING DATE: EARLIER FILING DATE: 1997-12-11  
 ; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/069, 750  
 ; PRIOR FILING DATE: EARLIER FILING DATE: 1997-12-16  
 ; SEQ ID NO 24  
 ; NUMBER OF SEQ ID NCS: 24  
 ; SOFTWARE: PatentIn Ver. 2.0  
 ; LENGTH: 1343  
 ; TYPE: DNA  
 ; ORGANISM: Porcine Circovirus Type II

Query Match 88.4%; Score 620.4; DB 10; Length 1768;  
 Best Local Similarity 92.7%; Pred. No. 2.3e-195;  
 Matches 651; Conservative 0; Mismatches 51; Indels 0; Gaps 0;

1 ATGACGTATCCAAGGAGGGCTTACCGAAGAGAACCCGGCAGGCATCTTGGC 60  
 |||||||

Db 1615 AAAATGGCATCTTCAACACCCGCCTCTGGGGACATGAGATTCAATTAATGACTGTAAGCGTAC 1556  
 Qy 181 ACAGTCAGAACGGCCCTCTGGGGACATGAGATTCAATTAATGACTTTCCT 240  
 Db 1555 ACAGTCACACGGCCCTCTGGGGGGACATGAGATTCAATTAATGACTTTCCT 1496  
 Qy 241 CCCCAAGGGGGCTCAAACCCCCGCTCTGTGGGGACATACAGAAATAAGAAG 300  
 Db 1495 GTTAAGGGTGAATTCTGGCCCTGCTGGGGGGACCAAAATCTCTATAACCCCTTGAATACTACAGAAATAAGAAG 1436  
 Qy 301 GTTAAGGGTGAATTCTGGCCCTGCTGGGGGGACATGAGATTCAATTAATGACTTTCCT 240  
 Db 1495 CCCGGGAGGGGGACAAATACTCTATAACCCCTTGAATACTACAGAAATAAGAAG 1436  
 Qy 361 AGTGCTGTTAGATGATAACTTTGTAAACAAAGGCCACAGGGCTCACCTATGACCCC 420  
 Db 1375 ACTGCTGTTAGATGATAACTTTGTAAACAAAGGCCACAGGGCTCACCTATGACCCA 1316  
 Qy 421 TATGTAACACTACTCTCCGGCATACCCAGCCCTCTACCAACTCCGGTAC 480  
 Db 1315 TATGTAACACTACTCTCCGGCATACCAACTCCGGTACCTACCCCTCCGGTAC 1256  
 Qy 481 TTACCCCCAACCTGTCTAGATTGATAACTTCCAACCAAACAAAGA 540  
 Db 1255 TTACACACCCAAACCTGTCTTGAACCTCCACATTGATAACTTCCAACCAAACAAAGG 1196  
 Qy 541 AACAGGCTGGCTGAGACTACAAACTGCTGGAAATGAGACCACGTAGGGCTCGGCACT 600  
 Db 1195 AATCAGCTTGGCTGAGGCTACAAACCTCTGGAAATGTGGACCACTAGGGCTCGGCACT 1136  
 Qy 601 GCGTTGAAACAGTATAACGACCAACTACAAATACCGTGTAACTGTACAA 660  
 Db 1135 GCGTTGAAACAGTAAATACGACCAACTACAAATACCGTGTAACTGTACAA 1076  
 Qy 661 TTACAGAGAATTAAATTAAAGGACCCCCACTTAACCCCTAA 702  
 Db 1075 TTACAGAGAATTAAATCTTAACCCACTTAACCCCTAA 1034

Db 1735 ATGACGTATCCAAGGAGGGCTTACCGAAGAGAACCCGGCAGGCATCTTGGC 1676  
 Qy 61 CAGATCCCTGGCGCCCTGGCTCACCCTGGCACCGTTACCGCTGGAGAAGG 120  
 Db 1675 CAGATCCCTGGCGCCCTGGCTCACCCTGGCACCGTTACCGCTGGAGAAGG 1616  
 Qy 121 AAAATGGCATCTCAACACCCGCCTCTCCGGCACCTCGGATATACTGTCAAGCGAAC 180  
 Db 1615 AAAATGGCATCTCAACACCCGCCTCTCCGGCACCTCGGATATACTGTCAAGCGAAC 1556  
 Qy 181 ACAGTCAGAACGGCCCTCTGGGGGGACATGAGATTCAATTAATGACTTTCCT 240  
 Db 1555 ACAGTCACACGGCCCTCTGGGGGGACCAAAATCTCTATAACCCCTTGAATACTACAGAAATAAGAAG 1436  
 Qy 241 CCCCAAGGGGGCTCAAACCCCCGCTCTGGGGGGACATGAGATTCAATTAATGACTTTCCT 240  
 Db 1495 CCCGGGAGGGGGACAAATACTCTATAACCCCTTGAATACTACAGAAATAAGAAG 1436  
 Qy 301 GTTAAGGGTGAATTCTGGCCCTGCTGGGGGGACAGGGGTACAGGGGACTGGGCTTC 360  
 Db 1435 GTTAAGGGTGAATTCTGGCCCTGCTGGGGGGACCAAAATACTCTATAACCCCTTGAATACTACAGAAATAAGAAG 1376  
 Qy 361 AGTGCTGTTAGATGATAACTTTGTAAACAAAGGCCACAGGGCTCACCTATGACCCC 420  
 Db 1375 ACTGCTGTTAGATGATAACTTTGTAAACAAAGGCCACAGGGCTCACCTATGACCCA 1316  
 Qy 421 TATGTAACACTACTCTCCGGCATACCCAGCCCTCTACCAACTCCGGTAC 480  
 Db 1315 TATGTAACACTACTCTCCGGCATACCAACTCCGGTACCTACCCCTCCGGTAC 1256  
 Qy 481 TTACCCCCAACCTGTCTAGATTGATAACTTCCAACCAAACAAAGA 540  
 Db 1255 TTACACACCCAAACCTGTCTTGAACCTCCACATTGATAACTTCCAACCAAACAAAGG 1196  
 Qy 541 AACAGGCTGGCTGAGACTACAAACTGCTGGAAATGAGACCACGTAGGGCTCGGCACT 600  
 Db 1195 AATCAGCTTGGCTGAGGCTACAAACCTCTGGAAATGTGGACCACTAGGGCTCGGCACT 1136  
 Qy 601 GCGTTGAAACAGTATAACGACCAACTACAAATACCGTGTAACTGTACAA 660  
 Db 1135 GCGTTGAAACAGTAAATACGACCAACTACAAATACCGTGTAACTGTACAA 1076  
 Qy 661 TTACAGAGAATTAAATTAAAGGACCCCCACTTAACCCCTAA 702  
 Db 1075 TTACAGAGAATTAAATCTTAACCCACTTAACCCCTAA 1034

Db 1135 GCCTTCGAAACAGTAAATACGACCAACTACAAATACCGTGTAACTGTACAA 1076  
 Qy 661 TTACAGAGAATTAAATTAAAGGACCCCCACTTAACCCCTAA 702  
 Db 1075 TTACAGAGAATTAAAGGACCCCCACTGTAAACCTAA 1034

## ALIGNMENTS



US-09-935-428A-24 ; LENGTH: 1768  
; TYPE: DNA  
; ORGANISM: Porcine circovirus  
; US-10-112-540-1

Query	Match	Score	DB	Length
QY	1 ATGACGTATCCAAGGAGGGTТАCCGAAGAAGAACCCGCCACGGCATCTTGGC	88.1%	10	1343;
Best Local Similarity	92.6%	Pred. No.	6.6e-195;	
Matches	650;	Mismatches	0;	
Indels	0;	Gaps	0;	
Db	1310 ATGACGTATCCAAGGAGGGTТАCCGCAAGAACCCGCCACGGCATCTTGGC	60		
QY	61 CAGATCCCGCCGCCCTGGCTCGTCCACCCCCGCCACCGTTACCGCTGGAGAAGG	120		
Db	1250 CAGATCCCGCCGCCCTGGCTCGTCCACCCCCGCCACCGCTACCGTTGGAGAAGG	1251		
QY	121 AAAATGGCATCTTCAACACCCGCCCTGGCATATTAAATGACTTTC	180		
Db	1190 AAAATGGCATCTTCAACACCCGCCCTGGCATATTAACTGTCAAGGCTAAC	1131		
QY	181 ACAGTCAGAACGGCCCTGGGACATGATGAGATTCAAAATATTGACTTTCTT	240		
Db	1130 ACAGTCAGAACGGCCCTGGGACATGATGAGATTAAATGACGACTTTGTT	1071		
QY	241 CCCCCAGGGGGGTCAAAACCCCGCTCTGTGCCCCTTGAAATFACTACAGAATAAGAAAG	300		
Db	1070 CCCCCGGAGGGGGGACCAACAAAATCTCTATACCCCTGGGAGATGAGATTAAATGACGACTTTGTT	1011		
QY	301 GTTAAGGGTTGAAATTAGTATAACTTGTAACTAGAATAAGAAAG	360		
Db	950 ACTGCTGTTATTCTAGTATAACTTGTAACTAGGCTTGTGGGGATAGGGGCTCC	891		
QY	421 TATGTAAACTCTCCGCCATACCATAAACCCAGCCCTCACCTATGACCCC	420		
Db	890 TATGTAAACTACTCTCCGCCATACATCCCCAACCTCTCCCTACCCGTTAC	831		
QY	481 TTTACCCCCAAACCTGTCCCTAGATTTCACATTATTGATTAACCTTCCAAACAAAGA	540		
Db	830 TTACACCCAAACCTGTCTTGTACCTTCCACATTGATTAACCAATAACAAAGG	771		
QY	541 AACCAGCTGTGGCTGAGACTACAAACTGTGGAAATGTAGACCTAGCTGGCACT	600		
Db	770 AATCAGCTTGGCTGAGCTACAAACCTCTAGAAATGTGGACCACTGGCACT	711		
QY	601 GCGTTCGAAAACAGTATATACGAGGAAATACAATATCCGTGTAACCCTAA	660		
Db	710 GCGTTCGAAAACAGTAAATACGACCCAGGACTACAATATCCGTGTAACCCTAA	651		
QY	661 TTCAAGAAATTAAATTAAAGACCCCCACTTAACCTTAA	702		
Db	650 TTCAAGAAATTAACTTAAAGACCCCCACTTAACCTTAA	609		

RESULT 4  
US-10-112-540-1/c  
; Sequence 1, Application US/10112540  
; Patent No. US2002017216A1  
; GENERAL INFORMATION:  
; APPLICANT: Liu, Qiang  
; APPLICANT: Tikoo, Suresh K.  
; APPLICANT: Willson, Philip  
; APPLICANT: Babiu, Lorne A.  
; TITLE OF INVENTION: METHODS TO CULTURE CIRCOVIRUS  
; FILE REFERENCE: 293102003100  
; CURRENT APPLICATION NUMBER: US/10/112,540  
; CURRENT FILING DATE: 2002-03-27  
; PRIOR APPLICATION NUMBER: US 60/279,173  
; PRIOR FILING DATE: 2001-03-27  
; NUMBER OF SEQ ID NOS: 3  
; SOFTWARE: FastSEQ for Windows Version 4.0  
; SEQ ID NO 1

RESULT 5  
US-09-784-962-1/c  
; Sequence 1, Application US/09784962  
; Patent No. US2002014631A1  
; GENERAL INFORMATION:  
; APPLICANT: Allan, Gordon M.  
; APPLICANT: Meehan, Brian M.  
; APPLICANT: Ellis, John A.  
; APPLICANT: Krakowka, George S.  
; APPLICANT: Audonne, Jean-Christophe F.  
; TITLE OF INVENTION: PORCINE CIRCOVIRUS AND PARVOVIRUS VACCINE  
; FILE REFERENCE: 454313-2338  
; CURRENT APPLICATION NUMBER: US/09/784,962  
; CURRENT FILING DATE: 2001-02-16  
; PRIOR APPLICATION NUMBER: 09/347,594

RESULT 10  
 US-09-784-962-4/C  
 Sequence 4, Application US/09784962  
 Patent No. US20020146431A1  
 GENERAL INFORMATION:  
 APPLICANT: ALLAN, Gordon M.  
 APPLICANT: MEEHAN, Brian M.  
 APPLICANT: ELLIS, John A.  
 APPLICANT: KRAKOWKA, George S.  
 APPLICANT: AUDONNET, Jean-Christophe F.  
 TITLE OF INVENTION: PORCINE CIRCOVIRUS AND PARVOVIRUS VACCINE  
 FILE REFERENCE: 454313-2338  
 CURRENT APPLICATION NUMBER: US/09/784, 962  
 CURRENT FILING DATE: 2001-02-16  
 PRIOR APPLICATION NUMBER: 09/347, 594  
 PRIOR FILING DATE: 1999-07-04  
 PRIOR APPLICATION NUMBER: 98 08777  
 PRIOR FILING DATE: 1998-07-06  
 NUMBER OF SEQ ID NOS: 5  
 SOFTWARE: PatentIn Ver. 2.1  
 SEQ ID NO 4  
 LENGTH: 1768  
 TYPE: DNA  
 ORGANISM: Porcine circovirus  
 US-09-784-962-4

Query Match 48.4%; Score 340; DB 10; Length 1768;  
 Best Local Similarity 92.3%; Pred. No. 3.5e-102;  
 Matches 358; Conservative 0; Mismatches 30; Indels 0; Gaps 0;

Qy 315 CTGGCCCTGCTCCCCGATCACCCAGGGTGAACAGGGAGTGGGCTCCAGTGCCTGTTATT 374  
 Db 1768 CTGGCCCTGCTCCCCATCACCCAGGGTGAACAGGGAGTGGGCTCCACTGCTGTTATTCT 1709

Qy 375 AGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGACCCCTATGAAACTACTC 434  
 Db 1708 AGATGATAACTTTGTAACAAAGGCCACAGCCCTAACCTATGAACTACTC 1649

Qy 435 CTCCCGCATACCATACCCAGGCCCTCTCCCTACCACTTACCCCTAACCTATGAACTACTC 434  
 Db 1648 CTCCCGCCATACCATACCCAGGCCCTCTCCCTACCACTTACCCCTAACCTATGAACTACTC 1589

Qy 495 TGTCTCTAGATTCACTATTGATTACTTCCAAACCAACAAAGAAACCCACTGTTGGCT 554  
 Db 1588 TGTCTCTGACTCCACTATTGATTACTTCCAAACCAATACCCCTAGAAATGAGCTTTGGCT 1529

Qy 555 GAGACTACAAACTGCTGGAAATGTAGACCACTGAGCTGGCCTCGGCACCTGCCTGGCT 614  
 Db 1528 GAGGCTACAAACCTCTAGAAATGAGCTGGACCACTGAGCTGGCCTCGGCACCTGCCTGGCT 1469

Qy 615 TATATAGACCGGAAATACAATATCCGGTAACTTACCCCTTAACCCCTAA 674  
 Db 1468 TCTTAAGGACCCCCACTTAACCCCTAA 1381

RESULT 12  
 US-09-884-514-4/C  
 Sequence 4, Application US/09884514  
 Patent No. US20020146432A1  
 GENERAL INFORMATION:  
 APPLICANT: ALLAN, Gordon  
 APPLICANT: MEEHAN, Brian  
 APPLICANT: CLARK, Edward  
 APPLICANT: HAINES, Deborah  
 APPLICANT: HASSARD, Lori  
 APPLICANT: HARDING, John  
 APPLICANT: CHARREYRE, Catherine E.  
 APPLICANT: CHAPPUIS, Gilles E.  
 APPLICANT: NEWTONARDS, Francis McNeilly  
 TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC  
 FILE REFERENCE: REAGENTS  
 CURRENT APPLICATION NUMBER: US/09/884, 514  
 CURRENT FILING DATE: 2000-06-19

RESULT 11  
 US-09-884-514-3/C  
 Sequence 3, Application US/09884514  
 Patent No. US20020146432A1  
 GENERAL INFORMATION:  
 APPLICANT: ALLAN, Gordon  
 APPLICANT: MEEHAN, Brian  
 APPLICANT: CLARK, Edward  
 APPLICANT: HAINES, Deborah  
 APPLICANT: HASSARD, Lori  
 APPLICANT: HARDING, John  
 APPLICANT: CHARREYRE, Catherine E.

1707 AGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGACCCCTATGTAACACTC 1648  
 4 35 CTCGGCCATACCATACCCAGCCCTTCTTACCAACTCCGGTACTTTACCCCCAAACC 494  
 1647 CTCGGCCATACCATACCCAGCCCTTCTTACCAACTCCGGTACTTTACCCCCAAACC 1588  
 4 95 TGTCCCTAGATTTCACTATTGATTACTTCCAACAAACAAAGAACAGCTGGCT 554  
 1587 TGTCCCTAGATTCCACTATTGATTACTTCCAACAAACAAAGAACATGGCTGGCT 1528  
 555 GAGACTACAAACTGCTGGAATGACCACGTAGGCTAGGCTAACCTTACCCCCAAACC 554  
 1527 GAGACTACAAACTACTGGAAATGAGACCACGTAGGCTAGGCTAACCTAACAG 1468  
 QY 615 TATATACGACCAGGAATACAAATATCCGGTGTAAACCATGTAATTCAAGAAATTAA 674  
 1467 TATATACGACCAGGAATACAAATATCCGGTGTAAACCATGTAATTCAAGAAATTAA 1408  
 Db 1587 TGTCCCTAGATTTCACTATTGATTACTTCCAACAAACAAAGAACAAACTAAC 1528  
 QY 555 GAGACTACAAACTGCTGGAATGACCACGTAGGCTAGGCTAACCTTACCCCCAAACC 614  
 Db 1527 GAGACTACAAACTACTGGAAATGAGACCACGTAGGCTAGGCTAACCTAACAG 1588  
 QY 615 TATATACGACCAGGAATACAAATATCCGGTGTAAACCATGTAATTCAAGAAATTAA 674  
 Db 1467 TATATACGACCAGGAATACAAATATCCGGTGTAAACCATGTAATTCAAGAAATTAA 1408  
 QY 675 TTTTAAAGACCCCCCACTTAACCTTAA 702  
 Db 1407 TCTTAAAGACCCCCCACTTAACCTTAA 1380  
 RESULT 9  
 US-09-784-962-3/c  
 ; Sequence 3, Application US/09784962  
 ; Patent No. US20020146431A1  
 ; GENERAL INFORMATION:  
 ; APPLICANT: ALLAN, Gordon M.  
 ; APPLICANT: MEEHAN, Brian M.  
 ; APPLICANT: ELLIS, John A.  
 ; APPLICANT: KRAKOWKA, George S.  
 ; APPLICANT: AUDONNET, Jean-Christophe F.  
 ; TITLE OF INVENTION: PORCINE CIRCOVIRUS AND PARVOVIRUS VACCINE  
 ; FILE REFERENCE: 454313-2338  
 ; CURRENT APPLICATION NUMBER: US/09/784, 962  
 ; CURRENT FILING DATE: 2001-02-16  
 ; PRIOR APPLICATION NUMBER: 09/347, 594  
 ; PRIOR FILING DATE: 1999-07-04  
 ; PRIOR APPLICATION NUMBER: 98 08777  
 ; PRIOR FILING DATE: 1998-07-06  
 ; NUMBER OF SEQ ID NOS: 5  
 ; SOFTWARE: PatentIn Ver. 2.1  
 ; SEQ ID NO 3  
 ; LENGTH: 1768  
 ; TYPE: DNA  
 ; ORGANISM: Porcine circovirus  
 US-09-784-962-3  
 Query Match 48.4%; Score 340; DB 10; Length 1768;  
 Best Local Similarity 92.3%; Pred. No. 3.5e-102;  
 Matches 358; Conservative 0; Mismatches 30; Indels 0; Gaps 0;  
 QY 315 CTGGCCCTGCTCCCCGATCACCCAGGGTGCACAGGGGACTGGCTCCAGTGCCTTATT 374  
 Db 1768 CTGGCCCTGCTCCCCCATCACCCAGGGTGTAGGGGACTGGCTCCACTGCTGTATTCT 1709  
 QY 375 AGATGATAACTTTGTAACAAAGGCCACAGGGCTTACCTATGAACTACTC 434  
 Db 1708 AGATGATAACTTTGTAACAAAGGCCACAGGGCTTACCTATGAACTACTC 1649  
 QY 435 CTCCCGCCATACCATACCCAGGCCCTACCTTCTCCCTACCCAAAC 494  
 Db 1648 CTCCCGCCATACCATACCCAAACCTTCTCCCTACCCAAAC 1589  
 Query Match 53.9%; Score 378.4; DB 10; Length 1767;  
 Best Local Similarity 98.5%; Pred. No. 6e-115;  
 Matches 382; Conservative 0; Mismatches 6; Indels 0; Gaps 0;  
 QY 315 CTGGCCCTGCTCCCCGATCACCCAGGGTGCAGGGGACTGGCTCCAGTGCCTTATT 374  
 Db 1767 CTGGCCCTGCTCCCCGATCACCCAGGGTGCAGGGGACTGGCTCCAGTGCCTTATTCT 1708  
 QY 375 AGATGATAACTTTGTAACAAAGGCCACAGGGCTTACCTATGAACTACTC 434  
 Db 1707 AGATGATAACTTTGTAACAAAGGCCACAGGGCTTACCTATGAACTACTC 1648  
 QY 435 CTGGCCCATACCATACCCAGGCCCTTCTCCCTACCCAAAC 494  
 Db 1647 CTGGCCCATACCATACCCAAACCTTCTCCCTACCCAAAC 1588  
 QY 495 TGTCCCTAGATTTCACTATTGATTACTTCCAAACAAACAAAGAACCTAAC 554  
 Db 1588 TGTCCCTGACTCCACTATTGATTACTTCCAAACAAACAAAGAACCTAAC 1529  
 QY 555 GAGACTACAAACTGCTGGAATGACCACGTAGGCTAGGCTAACCTAACAG 614  
 Db 1528 GAGGCTACAAACCTCTAGAAATGTTGACCTAACATGTAATTCAAGAAATTAA 1469  
 QY 615 TATATACGACCAGGAATACAAATATCCGGTGTAAACCATGTAATTCAAGAAATTAA 674  
 Db 1468 TATATACGACCAGGAATACAAATATCCGGTGTAAACCATGTAATTCAAGAAATTAA 1409  
 QY 675 TTTTAAAGACCCCCCACTTAACCTAAC 702  
 Db 1408 TCTTAAAGACCCCCCACTTAACCTAAC 1381

Best Local Matches	Similarity	Pred. No.	Score	DB	Length
470; Conservative	67.0%	1.7e-91;	273.4;	9;	7460;
0; Mismatches	0;				
222;	Indels	9;			
	Gaps	2;			
			38.9%	9;	7460;
			Best Local Matches	68.4%	
			Best Local Matches	68.4%	
			0; Mismatches	181;	
			Indels	9;	
			Gaps	2;	
			QY	1	ATGACGTATCCAAGGAGGGTACCGAAGAAGAACCCGGCAGCCATCTGGC
			60		60
			Db	1723	ATGACGTGGCCAAGGAGGGTACCGAAGAAGAACCCGGCAGCCATCTGG
			1664		1664
			QY	61	CAGATCCCTCGCCGCCCTGGCTCGTCCACCC - - - - -
			114		114
			Db	1663	AACATCCCTCGGAGAAGACCATATTGGACACACCCGGCTCTAGAGATGG
			1604		1604
			QY	115	AGAAGGAAAATGGCATCTTCAACACCCGGCTCTCCGACCCCTGGATAATAGTCAAG
			174		174
			Db	1603	CGCCGAAGAACGGGTATCTTCGAATTCGGCTTCTACAGAAATTGACTACCCATAAA
			1544		1544
			QY	175	CGAACACAGTCAGAACGGCCCTCTGGGGGACATGATGAGGATTCAGAAATTGAC
			234		234
			Db	1543	GGATACTCGCAGCCATCTTGAATTGGAAATCTACCTCAAACTAACATGGCCAG
			1487		1487
			QY	235	TTTCTTCCCCAGGGAGGGGGTCAAACCCGGCTCTGGGGGACATGATGAGGATTCAAAATTGAC
			294		294
			Db	1486	TTCCTCCCCCTCAAGGGGCCAACCCCTACCCCTACCTCAAACTAACATGGCTTT
			1427		1427
			QY	295	AGAAAGGTTAAGGTGAATTCTGGCCCTGCTCCCCGATCACCCAGGGTGACAGGGGAGTG
			354		354
			Db	1426	AGAAAGGCTAAATGAATTGGATCTGGTGTATCTGGATGCCAACCTTAATCAAAGAGGTGTT
			1367		1367
			QY	355	GGCTCCAGTGTGTATTAGTGTAAACTTTGATAACCTTGTAAACAAGGCCACGCCCTCACCTAT
			414		414
			Db	1366	GGTCCACTGTGTGTATCTGGATGCCAACCTTGTAAACCCCTCCACCAACTTGGCCTAT
			1307		1307
			QY	415	GACCCCTATGTAAACTACTCCGGCATACCCATAACCCAGCCCTCACCTCC
			474		474
			Db	1306	GACCCCTATATTAACTACTCCGGCATACCCAGCCCTCACCTACCCACTCC
			1247		1247
			QY	475	CGGTACTTTACCCCCAAACCTGTCTAGATTTCACATTGATTACTTCCACCAACAC
			534		534
			Db	1246	AGGTACTTCACCCCCAACCTGAGCTGGACCAAAACATTGATTGGTTCACCCAAATAAT
			1187		1187
			QY	535	AAAAGAACCCAGCTGTGGCTGAGACTACAAACTGCTGGAAATGTAGCCAGTAGGCCCTC
			594		594
			Db	1186	AAAAGAACCCAGCTGTGGCTCCATTAAATACCCACCAATGGCAGCCACAGGCCCTC
			1127		1127
			QY	595	GGCACTGGCTTCGAAACAGTATATAGCAGGAATAATTCGGTGTAAACCATGTAT
			654		654
			Db	1126	GGCTATGGCTCCAAAATGGCAGCCACAGGCCAAATTTAACCCACACCAATGTGGTAAGGCTGACTATTAT
			1067		1067
			QY	655	GTACAATTCAGAGAATTAAATTAAAGGCCCTTAACTTAA
			695		695
			Db	1066	GTACAATTCAGAGAATTAAATTAAAGGCCCTTAACTTAA
			1026		1026

Search completed: May 18, 2003, 13:00:07  
Job time : 170 secs

RESULT 15  
US-10-038-001-7/C  
Sequence 7, Application US/10038001  
Publication No. US2002018952A1  
GENERAL INFORMATION:  
APPLICANT: PALMER, Kenneth E.  
APPLICANT: POGUE, Gregory P.  
APPLICANT: MCCORMICK, Alison  
TITLE OF INVENTION: ROLLING CIRCLE REPLICON EXPRESSION  
TITLE OF INVENTION: VECTORS  
FILE REFERENCE: 008010179CPUS01  
CURRENT APPLICATION NUMBER: US/10/038, 001  
CURRENT FILING DATE: 2001-12-20  
PRIOR APPLICATION NUMBER: 09/505,477  
PRIOR FILING DATE: 2000-02-16  
NUMBER OF SEQ ID NOS: 9  
SOFTWARE: FastSEQ for Windows Version 4.0  
SEQ ID NO 7  
LENGTH: 7460  
TYPE: DNA  
ORGANISM: Porcine circovirus

; PRIOR APPLICATION NUMBER: FR 9800873  
 ; PRIOR FILING DATE: 1998-01-22  
 ; PRIOR APPLICATION NUMBER: FR 9803707  
 ; PRIOR FILING DATE: 1998-03-20  
 ; PRIOR APPLICATION NUMBER: FR 97/12382  
 ; PRIOR FILING DATE: 1997-10-03  
 ; NUMBER OF SEQ ID NOS: 6  
 ; SOFTWARE: PatentIn Ver. 2.0  
 ; SEQ ID NO: 4  
 ; LENGTH: 1768  
 ; TYPE: DNA  
 ; ORGANISM: Porcine circovirus  
 ; US-09-884-514-4

Query Match 48.48; Score 340; DB 10; Length 1768;  
 Best Local Similarity 92.38; Pred. No. 3.5e-102;  
 Matches 358; Conservative 0; Mismatches 30; Indels 0; Gaps 0;

Query	Match	Score	DB	Length
Qy	315 CTGGCCCTGCTCCCCGATCACCCAGGGTCCAGGGAGTGGCTTATT	339	10	1768
Db	1768 CTGGCCCTGCTCCCCCATCACCCAGGGTATAAGGGAGTGGCTTATT	375	1768	1709
Qy	375 AGATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGTAACACTC	374	1708	1649
Db	1708 AGATGATAACTTTGTAACAAAGGGTGTAGGGAGTGGCTTATTCT	434	1708	1649
Qy	435 CTCCGGCCATACCATAACCCAGGCCCTTCTCCTACCATATGTAACACTC	494	1648	1589
Db	1648 CTCCGGCCATACCATACCCCAACCCCTCTCCTACCATCCGGTACTTCA	554	1648	1589
Qy	495 TGTCTAGATTCACTATTGATTACTTGACCCCTTACCTATGTAACACTC	554	1588	1529
Db	1588 TGTTCTTGACTCCACTATTGATTACTTGACCCCTTACCTATGTAACACTC	614	1588	1529
Qy	555 GAGACTACAAACTGCTGGAAATGTAGACCAACGACTGCCCTCGGAAACAG	614	1528	1469
Db	1528 GAGACTACAAACCTCTGGAAATGTGGACCAACGACTGCCCTCGGCTCGGAAACAG	674	1528	1469
Qy	615 TATATAGACCAGGAATAACATATCCGGTAAACCATGATGTACAATTCAAGAGAATTAA	674	1468	1409
Db	1468 TAAATAGACCAGGAATAACATATCCGGTAAACCATGATGTACAATTCAAGAGAATTAA	702	1468	1409
Qy	675 TTTAAAGACCCCCCACTTAACCCCTTAA	702	1408	1381
Db	1408 TCTAAAGACCCCCCACTTAACCCCTTAA	702	1408	1381

RESULT 13 US-09-884-514-6/c

; Sequence 6, Application US/09884514  
 ; Patent No. US20020146432A1  
 ; GENERAL INFORMATION:  
 ; APPLICANT: ALLAN, Gordon  
 ; APPLICANT: MEEHAN, Brian  
 ; APPLICANT: CLARK, Edward  
 ; APPLICANT: HAINES, Deborah  
 ; APPLICANT: HARDING, John  
 ; APPLICANT: CHARREYRE, Catherine E.  
 ; APPLICANT: CHAPUIS, Gilles E.  
 ; APPLICANT: NEWTONARDS, Francis McNeilly  
 ; TITLE OF INVENTION: NEW PORCINE CIRCOVIRUSES, VACCINES AND DIAGNOSTIC  
 ; FILE REFERENCE: AILAN  
 ; CURRENT APPLICATION NUMBER: US/09/884,514  
 ; CURRENT FILING DATE: 2000-06-19  
 ; PRIOR APPLICATION NUMBER: FR 9800873  
 ; PRIOR FILING DATE: 1998-01-22  
 ; PRIOR APPLICATION NUMBER: FR 9803707  
 ; PRIOR FILING DATE: 1998-03-20  
 ; PRIOR APPLICATION NUMBER: FR 97/12382  
 ; PRIOR FILING DATE: 1997-10-03  
 ; NUMBER OF SEQ ID NOS: 6

Query Match 43.88; Score 307.8; DB 10; Length 1759;

; SOFTWARE: PatentIn Ver. 2.0  
 ; SEQ ID NO: 6  
 ; LENGTH: 1759  
 ; TYPE: DNA  
 ; ORGANISM: Porcine Circovirus Type I  
 ; US-09-935-428A-2

Query Match 48.38; Score 339; DB 10; Length 1768;  
 Best Local Similarity 92.28; Pred. No. 7.5e-102;  
 Matches 357; Conservative 0; Mismatches 30; Indels 0; Gaps 0;

Query	Match	Score	DB	Length
Qy	316 TGGCCCTGCTCCCCGATCACCCAGGGTACAGGGAGTGGCTCCAGTGCTTATT	339	10	1768
Db	1768 TGGCCCTGCTCCCCCATCACCCAGGGTATAAGGGAGTGGCTCCAGTGCTTATT	375	1768	1709
Qy	376 GATGATAACTTTGTAACAAAGGCCACAGCCCTCACCTATGTAACACTC	435	1708	1649
Db	1708 GATGATAACTTTGTAACAAAGGCCACAGCCCTAACCTATGTAACACTC	435	1708	1649
Qy	436 TCCCGCCATACCCATAACCCCTTCTCCCTACACTCCAAACCT	495	1648	1589
Db	1648 TCCCGCCATACCCATAACCCCTTCTCCCTACACTCCAAACCT	555	1648	1589
Qy	496 GTCTAGATTCACTATTGATTACTTCAACCAACAAACAGCTGGGCTG	555	1588	1529
Db	1588 GTTCTGACTCCACTATTGATTACTTCAACCAAAAGGAATCAGCTTGGCTG	615	1588	1529
Qy	556 AGACTACAAACTGCTGGAAATGTAGACCACTAGGCCACTGGCTTGGAAACAGT	615	1528	1469
Db	1528 AGGCTACAAACCTCTGAATACTCCCTCTGAATACTCCCTAGGCTTGGAAACAGT	616	1528	1469
Qy	616 ATATAGCCAGGAATAACATATCCGGTGTACCATGATGTACATTAA	675	1468	1409
Db	1468 ATATAGCCAGGAATAATCCGGTGTACCATGATGTACATTCAAGAAATTAA	702	1468	1409

RESULT 14 US-09-935-428A-2/c

; Sequence 2, Application US/09935428A  
 ; Patent No. US20020106639A1  
 ; GENERAL INFORMATION:  
 ; APPLICANT: WANG, Li  
 ; APPLICANT: BABIUK, LORNE A.  
 ; APPLICANT: POTTER, ANDREW A.  
 ; APPLICANT: WILLSON, PHILIP  
 ; TITLE OF INVENTION: POSTWEANING MULTISYSTEM WASTING SYNDROME VIRUS FROM PIGS  
 ; FILE REFERENCE: 9000-0040  
 ; CURRENT APPLICATION NUMBER: US/09/935,428A  
 ; CURRENT FILING DATE: 2001-08-20  
 ; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/209, 961  
 ; PRIOR FILING DATE: EARLIER FILING DATE: 1998-12-10  
 ; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/069, 233  
 ; PRIOR FILING DATE: EARLIER FILING DATE: 1997-12-11  
 ; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/069, 750  
 ; PRIOR FILING DATE: EARLIER FILING DATE: 1997-12-16  
 ; NUMBER OF SEQ ID NOS: 24  
 ; SOFTWARE: PatentIn Ver. 2.0  
 ; SEQ ID NO: 2  
 ; LENGTH: 1759  
 ; TYPE: DNA  
 ; ORGANISM: Porcine Circovirus Type I  
 ; US-09-935-428A-2